Table 1-1

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						MPM-117	MPM-118/	MPM-119 /	Tumor Aver
_		Tumors	BPH		Aver. BPH		Aver. BPH		1
	788180	25.18	0.50	102.19	5.44	15.38	90.33	40.12	50.69
	308682	8.99	0.27	26.49	39.49	22.07	29.80	46.70	32.91
	646037	24.42	0.75	53.07	14.65	28.67	17.92	49.22	32.70
	281003	128.77	4.90	33.20		7.16			26.26
	1034473	33.61	1.35	44.07		18.38	21.16		24.97
	284701	33.05	1.53	•		11.69			21.68
	66336	1.67	0.10			26.08			1
	281039	14.05	0.90	•		1.52			i .
	26736	9.52	l .	1		3.08			1
	784168	53.04	1	1		16.92			13.84
	486035	24.75	1	1					ž .
	593185	28.06	t .	I .					I .
	222157	8.80		i .					1
	35058	15.59	1						li .
	730739		1	1					1
	1636908	t .	1						1
	292515	i	3	1					*
	450049		1	T .					
-3	666334		ı						•
	32160		1						•
4	26519	1	l .	1					1
	42415	•					1		
I	50772								1
	1412481	1	1	1					i i
	197525	1		1					i
i.	490178	1	3	1					
	461522	1	3	1					
	746152	1		1					1
::Ē	71295	1	1	1					1
U	81405	1	1	1					
	161044		1	l l			48 9.0		
	82585							36 5.1	1
ajb	59529				22 5.7				80 8.16
	59569	•	•	96 26.			••	-	95 8.08
	48896 25937				23 5.3		68 15.°		79 8.04
	28208	I		3	99 11.5		50 23.		88 7.92
	82394	•		1		80 21.			77 7.91
	44771	1		88 - 7.					00 7.90
	104881				.33 12.				52 7.78
	4610				.53 17.				20 7.69
	4602	3							.24 7.59
	68729			4					.95 7.48
	7573	ı.		1					.46 7.41
	7312		E .	1					.10 7.40
	15565	1							.24 7.33
	8261	•		4					.60 7.27
	8257		1						.04 7.26
	221			l l					5.71 7.24
				•					-

283173	17.85	2.47	17.57	3.98	4.39	7.32	2.81	7.21
825822	14.07	1.97	5.87	8.14	6.53	10.54	4.71	7.16
280217	20.52	2.89	9.65	0.93	2.71	20.73	1.45	7.09
262053	10.75	1.52	5.67	9.07	8.71	8.34	3.60	7.08
66317	5.50	0.78	15.41	4.56	4.00	10.34	0.99	7.06
565235	35.83	5.15	3.19	8.34	3.79	17.38	2.08	6.96
1468260	4.53	0.65	13.06	1.36	7.45	4.03	8.89	6.96
826103	1.31	0.19	8.19	7.96	5.94	9.88	2.70	6.93
796366	34.94	5.05	10.57	5.34	12.35	3.13	3.18	6.92
877835	23.06	3.34	9.71	3.97	5.61	7.41	7.82	6.90
258300	18.15	2.65	4.31	7.13	13.32	1.26	8.27	6.86
814062	0.81	0.12	10.04	5.59	8.22	8.85	1.42	6.83
1034738	22.40	3.29	5.92	5.75	6.61	9.70	6.12	6.82
951313	6.84	1.01	2.78	4.19	15.48	7.44	3.86	6.75
253132	1.40	0.21	8.70	9.55	4.09	5.72	5.23	6.66
824510	1.38	0.21	8.56	9.50	7.92	5.67	1.59	6.65
826072	1.20	0.18	7.01	8.64	4.02	10.85	2.42	6.59
796754	0.66	0.10	6.20	5.19	5.58	9.09	6.47	6.50
251806	2.94	0.46	6.74	6.71	5.58	9.16	3.74	6.39
840306	2.49	0.39	2.46	5.82	8.43	9.48	5.73	6.38
815163	1.20	0.19	6.07	6.53	9.54	7.44	2.20	6.36
838285	1.34	0.21	4.71	7.31	6.15	8.05	5.54	6.35
565863	3.93	0.63	13.52	9.05	2.58	3.19	3.01	6.27
757435	365.78	58.35	3.90	3.17	17.54	3.19	3.55	6.27
67401	1.47	0.23	1.08	1.27	26.14	1.90	0.96	6.27
796239	26.07	4.18	1.88	2.16	10.01	7.23	9.94	6.24
321706	7.80	1.25	8.36	1.45	14.65	1.76	4.95	6.23
564801	12.35	1.99	5.89	2.75	17.66	3.44	1.34	6.21
950778	3.00	0.49	12.47	7.19	4.14	3.62	3.50	6.18
884606	33.46	5.43	6.01	3.92	16.90	2.25	1.74	6.16
454317	20.56	3.34	3.52	4.38	7.94	12.41	2.55	6.16
796646	6.21	1.01	5.13	3.61	5.34	6.22	10.48	6.16
840894	1 1	5.66	2.34	6.76	5.31	9.85	6.47	6.15
52933	1 1	0.58	5.96	6.27	4.93	6.70	6.87	6.14
594994		2.38	5.57	4.29	8.11	6.75	5.98	6.14
586780		0.31	11.12	7.73	4.39	3.68	3.68	6.12
868282		1.47	5.69	6.02	6.96	6.08	5.84	6.12 6.00
1535590		66.91	5.47	9.25	6.43	4.48	4.39	5.99
25997		0.28	6.61	6.24	5.06	6.62	5.42	5.98
83908			10.16	10.63	1.88	5.07	2.14	5.97
35966			4.04	9.90	3.64	7.90	4.38	5.95
50973			4.92	3.91	8.19	7.51	5.20 2.17	5.94
77006	t t		1.49	2.31	5.43	18.27	5.38	5.92
15105	1		5.93	10.61	2.93	4.74	4.59	5.88
84292		1	2.96	5.54	10.03	6.26 6.06	1.00	5.87
82412		1 1	8.58	7.60	6.12		1	5.84
81466		1 1	5.53	5.17	7.24	9.88	1.37 1.99	5.84 5.84
105617	•	1 1	6.48	4.11	1.30	15.30	0.84	5.83
84094	l l		11.87	1.67	12.92	1.88 4.59	5.11	5.78
8115		1 1	8.01	4.53	6.69	4.39 9.32	3.03	5.71
16372	82 4.01	0.70	2.86	2.47	10.87	7.32	3.03	J./1

eseesal	1.00	0.221	C 10	E 0E	4.20	7.63	4 22	5.71
838807 77915	1.90 14.99	0.33 2.63	6.10 3.34	5.85 8.82	4.20 3.66	7.62 2.66	4.77 10.02	5.71 5.70
252314	0.69	0.12	3.34 8.50	5.50	5.00 6.04	2.00 3.55	4.72	5.66
814235	12.37	2.19	6.17	6.36	5.75	7.43	2.52	5.65
461476	0.78	0.14	7.32	7.96	5.75 5.15	7.43 5.06	2.71	5.64
1		2.26	4.29	7. 90 5.26	8.83	2.50	7.34	5.64
754358	12.77	0.24	4.29 3.38	5.20 6.17	9.26	3.91	5.41	5.63
53022 859228	85.06	15.20	5.32	4.79	9.20 6.57	7.30	4.01	5.60
290370	0.62	0.11	3.32 4.96	4.7 9 3.44	7.59	8.28	3.41	5.54
81331	2.84	0.52	1.67	4.49	1.34	18.34	1.80	5.53
260116	0.88	0.16	6.00	5.67	4.91	5.69	5.33	5.52
416374	4.49	0.10	3.17	8.84	3.61	10.29	1.67	5.52
840606	15.85	2.88	3.74	4.47	5.11	9.38	4.79	5.50
51773	11.48	2.09	6.89	4.08	3.70	9.38 8.77	4.03	5.50
240694	4.53	0.83	5.08	8.53	3.38	3.73	6.64	5.47
1605407	9.00	1.65	4.85	3.69	10.03	6.25	2.42	5.45
139771	7.72	1.42	13.10	8.46	1.72	1.70	2.24	5.44
1630990	20.64	3.80	8.87	4.51	3.52	9.31	0.98	5.44
1390584	6.54	1.21	3.13	4.93	5.15	12.83	0.99	5.41
186918	3.04	0.56	2.90	5.49	4.54	11.24	2.73	5.38
462097	8.32	1.55	7.30	2.88	1.25	9.95	5.50	5.38
32381	2.18	0.41	1.75	2.88	9.49	3.19	9.57	5.38
491764	0.81	0.15	6.00	5.07	5.46	5.33	4.98	5.37
1456160	41.61	7.76	2.23	3.67	7.39	11.23	2.29	5.36
159608	14.43	2.70	10.47	1.64	1.63	7.83	5.21	5.36
80050	8.15	1.52	6.41	4.16	11.93	0.95	3.31	5.35
610124	1.70	0.32	4.17	6.14	5.42	5.63	5.22	5.32
280763	0.53	0.10	3.60	5.23	5.40	6.52	5.64	5.28
283312	63.94	12.14	6.26	3.00	2.86	6.54	7.69	5.27
359781	7.82	1.49	3.10	9.11	3.52	6.46	4.12	5.26
297439	48.60	9.26	5.11	9.34	4.69	2.38	4.73	5.25
731410	0.53	0.10	5.74	4.50	5.23	5.26	5.48	5.24
772416	1.03	0.20	4.21	4.42	4.71	7.57	5.02	5.19
267419	0.97	0.19	3.98	9.05	3.92	5.50	3.38	5.16
897807	5.08	0.99	5.07	5.96	9.08	3.03	2.65	5.16
593658	1.08	0.22	4.01	5.03	5.25	5.65	5.29	5.04
511952	225.66	45.17	0.81	4.98	1.25	15.74	2.20	5.00
743452	7.90	1.59	7.90	7.46	2.03	4.53	2.98	4.98
878281	6.80	1.37	5.03	1.94	7.14	8.50	2.27	4.98
825297	0.96		4.48	3.92	7.31	4.87	4.28	4.97
731016			6.76	6.19	3.31	4.49	4.06	4.96
824723			3.64	3.92	5.23	8.15	3.87	4.96
824122			6.80	5.17	5.77	4.87	2.17	4.95
755239	1		5.25	3.08	3.43	8.49	4.50	4.95
784130	t .		3.20	5.09	7.85	4.99	3.50	4.93
1585549	B		5.67	6.28	3.85	4.93	3.89	4.92
701677	1		7.12	3.55	3.14	8.18	2.54	4.91
814017	1	3 1	6.83	7.64	4.06	2.12	3.86	4.90
530185			2.68	5.31	6.79	3.57	6.04	4.88
773283			4.28	4.12	3.43	7.01	5.50	4.87
19472	5.22	1.07	8.66	7.32	2.79	2.88	2.66	4.86

788655	7.66	1.58	4.19	3.33	7.19	6.38	3.22	4.86
238661	2.74	0.56	9.41	7.84	2.13	2.69	2.21	4.86
773248	21.90	4.51	1.37	2.18	6.84	6.57	7.31	4.86
124252	6.06	1.25	11.29	6.83	1.86	2.10	2.21	4.85
814306	18.28	3.77	3.29	5.66	3.89	7.34	4.08	4.85
754280	23.83	4.92	9.79	5.07	5.33	1.97	2.08	4.85
1636108	3.58	0.74	5.59	4.80	5.19	4.56	4.09	4.85
461592	11.87	2.45	8.03	6.34	4.05	3.94	1.87	4.84
771053	16.14	3.34	3.85	4.29	4.79	3.56	7.67	4.83
703943	4.77	0.99	3.01	6.34	6.79	4.06	3.91	4.82
823691	7.42	1.54	2.71	4.70	4.41	8.34	3.93	4.82
289071	37.48	7.78	3.68	4.93	11.18	2.30	2.00	4.82
510576	132.22	27.52	5.44	2.70	2.95	8.26	4.68	4.81
264465	0.77	0.16	5.25	4.74	5.40	3.85	4.73	4.79
511428	24.31	5.08	5.64	3.21	4.36	5.17	5.56	4.79
488431	10.48	2.19	6.83	4.34	3.39	2.92	6.45	4.78
359701	5.21	1.09	4.50	5.09	3.54	3.24	7.52	4.78
1606837	23.76	4.98	6.95	2.67	3.34	5.80	5.10	4.77
823615	9.40	1.97	6.36	2.61	10.35	2.83	1.67	4.76
795265	4.58	0.97	1.65	3.44	3.71	12.61	2.30	4.74
813275	1.75	0.37	2.37	3.77	3.58	8.89	5.04	4.73
796351	2.34	0.50	5.27	6.47	3.71	4.50	3.66	4.72
787938	5.09	1.08	1,71	5.35	8.20	6.29	2.02	4.71
375682	31.04	6.60	3.43	4.82	7.07	3.67	4.53	4.70
869450	51.58	10.98	10.12	2.49	0.92	4.42	5.54	4.70
251435		0.16	4.68	5.34	5.04	4.18	4.15	4.68
824267		0.16	4.49	4.65	4.35	4.86	5.03	4.68
214006		1.00	9.34	7.13	2.17	3.26	1.38	4.66
251517		0.16	4.87	4.91	5.29	3.94	4.18	4.64
609052		0.19	5.08	4.85	3.84	4.06	5.34	4.63
1466631		0.26	3.47	5.67	4.78	4.73	4.50	4.63
1500000		1.32	8.25	6.21	3.19	2.69	2.81	4.63
1055201	1 1	6.55	4.44	2.37	4.47	9.42	2.38	4.62
840878	1	5.59	2.71	4.00	3.35	9.63	3.38	4.62
773204	B		2.70	3.84	5.75	1.24	9.53	4.61
290841	1	1 1	7.54	6.98	2.99	3.39	2.13	4.61
134969	6.81	1.49	10.24	6.19	2.21	2.17	2.11	4.58
824376	0.49	0.11	5.18	4.11	6.01	6.67	0.94	4.58
768643	41.57	9.10	2.42	4.02	11.03	2.91	2.45	4.57
2661	7 4.62	1.01	1.74	7.29	3.24	7.49	3.02	4.56
30553	8 11.44	2.54	6.22	5.46	2.58	2.41	5.84	4.50
81116	2 6.36	1.41	4.62	2.07	5.28	2.80	7.76	4.50
163734	3 5.03	1.12	3.26	4.54	4.14	6.66	3.91	4.50
19777	9 8.61	1.91	10.37	7.03	1.68	1.51	1.91	4.50
81554			6.84	5.83	3.95	4.46	1.38	4.49
5170		•	4.47	4.07	4.99	3.08	5.73	4.47
75720	1		3.96	4.49	6.00	3.91	3.98	4.47
83876			6.93	6.35	2.56	3.70	2.80	4.47
34398		1 1	2.47	4.92	6.23	2.51	6.15	4.46
76740		, ,	5.11	3.93	4.22	4.46	4.52	4.45
7903	37.5	8 8.53	2.88	8.31	5.01	3.70	2.12	4.41

815036	1.62	0.37	5.65	5.86	3.71	4.71	2.08	4.40
823895	1.57	0.36	4.08	4.25	4.38	4.82	4.45	4.40
201282	6.42	1.46	1.85	6.04	4.99	6.43	2.63	4.39
773208	26.40	6.05	2.45	3.37	5.47	1.22	9.32	4.37
121436	6.37	1.46	7.91	6.01	2.52	3.24	2.09	4.36
238821	6.54	1.50	5.59	1.57	4.76	0.92	8.90	4.35
814798	32.86	7.58	3.15	4.62	2.68	9.26	1.99	4.34
196650	10.93	2.52	4.17	6.60	2.20	6.71	1.99	4.33
971367	35.22	8.15	4.78	2.95	2.51	8.35	3.04	4.32
841220	1.46	0.34	3.52	5.00	4.19	4.54	4.37	4.32
200599	13.42	3.11	5.88	2.44	8.89	1.51	2.87	4.32
487733	15.01	3.48	3.75	5.14	4.04	6.05	2.61	4.32
564537	1.79	0.42	2.77	5.08	5.29	4.04	4.38	4.31
50606	14.94	3.47	5.20	2.99	5.04	3.92	4.39	4.31
253314	1.52	0.35	7.93	5.83	2.29	2.55	2.92	4.30
417908	12.10	2.82	8.40	4.93	3.98	1.81	2.34	4.29
29063	3.34	0.78	2.94	6.06	4.11	3.85	4.48	4.29
773345	1.21	0.28	3.42	4.56	2.89	8.45	2.12	4.29
1592006	5.30	1.24	4.98	3.66	4.06	4.48	4.22	4.28
461499	2.26	0.53	2.94	3.53	6.30	7.11	1.48	4.27
815800	1.22	0.29	5.36	6.04	5.43	3.31	1.13	4.25
436769	168.72	39.68	3.44	3.03	2.92	5.33	6.54	4.25
427838	4.05	0.95	7.99	6.36	2.29	2.32	2.29	4.25
50188	10.75	2.53	4.92	4.45	4.01	3.41	4.46	4.25 4.25
1519515	0.85	0.20	3.89	3.77	5.31	3.37	4.90	4.23 4.24
138444	4.04	0.95	3.84	7.52	3.07	4.21	2.54	4.2 4 4.24
1631849	25.01	5.91	4.16	5.55	3.42	4.66	3.40 4.94	4.24
613237	1.07	0.25	3.02	4.22	3.83	5.08 2.13	4.52	4.21
358850		5.06	7.24	3.98	3.18 3.92	3.90	5.86	4.20
290378		4.07	3.83	3.50	5.89	5.18	1.97	4.19
898109		1.62	4.54	3.37	3.89 4.26	6.62	2.60	4.18
82734		1.73	3.01	4.42	4.26 3.75	1.27	2.10	4.17
450307	1 .	2.25	7.70	6.02 2.52	9.06	1.63	1.24	4.17
593929	1 1	4.26	6.37	2.52 3.53	7.08	1.03	1.53	4.16
587415		3.12	7.46	5.06	5.56	2.68	3.67	4.16
842882			3.81 5.29	7.59	0.53	6.00	1.32	4.14
896914 1607765	3		3.29 4.88	3.54	3.13	5.17	3.98	4.14
824329	1		4.04	4.22	5.97	3.83	2.60	4.13
155841			0.59	1.24	9.68	5.69	3.37	4.11
453599			3.89	3.56	3.50	6.56	3.01	4.11
48653	T.		3.08	5.14	3.85	5.78	2.65	4.10
71221			5.35	4.81	4.78	4.53	1.03	4.10
75316	i		2.94	3.35	2.20	9.53	2.46	4.10
163729			4.17	2.86	2.82	6.06	4.57	4.09
89824	1		2.92	4.71	4.80	5.15	2.87	4.09
77104		1 1	5.50	3.59	3.73	3.20	4.39	4.08
95112		1 1	3.30	7.80	2.31	3.51	3.45	4.07
61115		1 1	4.60	2.92	4.15	5.04	3.66	4.07
84292	1	1 1	3.78	3.80	4.94	4.96	2.87	4.07
82484	1	1 1	5.28	3.89	3.11	6.61	1.41	4.06
J2 10	-1	11	3				•	

878544	19.20	4.73	8.54	2.29	4.85	2.59	2.02	4.06
781366	12.06	2.98	4.50	3.42	1.61	2.86	7.87	4.05
79000	6.24	1.54	2.16	5.95	3.81	5.31	3.04	4.05
24208	7.49	1.85	3.19	3.67	3.57	4.54	5.28	4.05
729953	4.95	1.23	3.26	4.43	3.91	6.32	2.25	4.04
34134	2.73	0.68	6.48	3.68	4.71	3.19	2.08	4.03
745283	7.94	1.97	4.03	3.25	3.72	4.68	4.44	4.02
1602619	135.32	33.71	4.29	3.43	3.41	4.71	4.23	4.02
1636868	5.07	1.27	5.31	4.50	3.81	4.48	1.96	4.01
726483	3.50	0.87	2.71	4.78	2.55	5.48	4.52	4.01
360403	28.16	7.03	9.60	1.99	5.91	1.37	1.17	4.01
757462	12.77	3.19	4.05	4.75	4.11	4.73	2.41	4.01
279482	8.66	2.16	3.13	2.00	10.92	1.42	2.54	4.00
811766	4.93	1.23	1.80	3.53	4.42	6.47	3.79	4.00
50764	0.43	0.11	4.10	3.54	3.73	5.17	3.48	4.00
626716	21.60	5.41	2.78	6.84	3.77	3.86	2.73	4.00
213682	13.07	3.27	0.79	3.49	2.68	9.05	3.96	3.99
685922	3.59	0.90	3.85	3.96	3.20	5.02	3.93	3.99
1631194	21.83	5.47	5.50	1.73	5.19	6.07	1.46	3.99
823578	0.56	0.14	4.98	3.43	4.18	3.36	3.93	3.98
32509	4.38	1.10	1.50	7.60	1.95	4.07	4.72	3.97
845352	13.36	3.37	4.44	3.86	3.95	2.54	5.05	3.97
191978		12.11	3.93	4.27	3.76	5.36	2.48	3.96
120707	1 I	4.95	2.32	3.87	3.65	8.32	1.62	3.96
75415	1 1	13.09	4.45	4.47	2.60	4.53	3.70	3.95
186768		2.39	2.81	1.73	11.34	2.18	1.69	3.95
80484	1 1	3.72	9.35	2.80	1.75	3.52	2.32	3.95
1475200	1 1	7.14	1.70	3.21	6.52	2.96	5.34	3.95
810510	1 1	7.03	3.76	3.64	2.68	4.64	5.00	3.94
897690		4.03	5.68	4.08	4.14	1.92	3.87	3.94
45327		2.08	2.60	3.75	4.04	5.11	4.18	3.94
299723	1	0.76	1.71	2.92	5.37	4.69	4.93	3.93
81578	12.77	3.25	5.02	3.62	6.26	2.36	2.37	3.93
19788	8 4.97	1.27	4.11	2.76	2.65	3.61	6.49	3.92
81057	1 4.37	1.11	2.29	5.21	2.51	6.03	3.57	3.92
76844	3 26.58	6.79	2.54	4.37	3.56	7.05	2.07	3.92
15031		1.35	2.49	4.79	3.04	6.51	2.76	3.92
19618	9 11.19	2.86	4.68	4.67	3.98	2.71	3.52	3.91
62699	0 7.39	1.89	2.81	2.77	3.38	7.85	2.73	3.91
12739	6 14.07		3.73	5.03	1.41	6.75	2.59	3.90
4971	0 3.21	1	2.93	3.48	4.65	4.08	4.34	3.90
26286	36.97		3.65	2.59	3.11	5.73	4.36	3.89
81573		1 1	3.91	4.25	3.43	5.08	2.74	3.88
25668	t t		3.22	3.59	3.08	4.91	4.60	3.88
46150			4.43	4.30	3.23	5.77	1.64	3.87 3.87
8095	3		5.73	2.71	3.10	4.48	3.32	3.87
8111	1	1	2.85	4.21	4.20	4.46	3.62	3.87
5470	1		4.03	2.83	3.49	4.83	4.11	3.86 3.86
10357			3.91	1.25	4.74	0.71	8.68	3.86 3.86
8247	4		7.31	4.22	2.34	3.52	1.89 3.95	3.85
8150	17 14.8	3.84	3.95	3.11	3.65	4.61	3.73	2.02

							4 601	2 05
32684	24.32	6.32	4.79	1.68	2.94	5.24	4.60	3.85
178137	238.41	61.93	3.66	2.98	2.56	6.38	3.66	3.85
282283	0.84	0.22	3.28	3.36	3.61	4.89	4.09	3.85
725978	4.92	1.28	5.17	3.07	2.63	4.05	4.31	3.85
1631762	46.32	12.06	2.78	4.97	2.53	4.94	4.00	3.84
1553306	7.92	2.06	3.60	3.69	3.79	4.48	3.64	3.84
590145	4.37	1.14	3.78	3.14	3.77	5.80	2.69	3.84
815230	9.59	2.50	3.37	6.27	3.72	3.68	2.14	3.84
824237	74.89	19.54	3.14	3.18	3.55	5.72	3.57	3.83
361048	3.13	0.82	2.53	3.04	6.30	4.44	2.84	3.83
417867	13.52	3.54	2.81	3.70	3.15	4.96	4.47	3.82
878152	4.20	1.10	3.49	3.90	4.76	3.69	3.25	3.82
487499	4.41	1.16	2.68	3.09	8.83	3.10	1.37	3.82
712570	1.32	0.35	4.30	4.75	4.52	3.37	2.10	3.81
124824	30.85	8.13	3.79	2.43	2.17	3.37	7.23	3.80
626385	15.94	4.20	2.97	4.37	4.40	2.13	5.11	3.79
813271	0.96	0.25	3.16	4.24	3.70	4.31	3.56	3.79
561971	3.81	1.01	2.50	4.39	4.74	3.83	3.46	3.79
824358	0.96	0.25	4.80	3.84	3.71	5.67	0.90	3.79
741067	64.30	17.00	0.88	4.25	2.30	7.09	4.38	3.78
46411	3.07	0.81	2.37	2.25	5.61	3.01	5.66	3.78
594517	24.81	6.57	4.24	1.83	7.82	1.45	3.53	3.78
825668		0.60	3.80	3.91	4.97	3.99	2.20	3.77
362926	4.39	1.17	1.58	4.23	3.24	2.08	7.71	3.77
814018	0.98	0.26	3.30	3.43	6.07	5.33	0.71	3.77
745503	42.62	11.32	2.92	3.95	3.33	3.03	5.59	3.77
884531	7.71	2.05	2.78	3.56	3.60	6.93	1.94	3.76
210862	2.56	0.68	5.33	3.36	3.68	3.66	2.76	3.76
504544	12.31	3.28	2.35	4.09	2.54	7.86	1.95	3.76
838668		0.95	1.88	5.28	2.47	7.20	1.95	3.76
433253		1.95	2.88	2.93	2.56	6.57	3.78	3.74
461330		1 1	3.19	3.49	3.78	5.20	3.05	3.74
87852			4.03	4.61	3.75	2.79	3.49	3.74
38318			6.71	2.73	7.18	1.05	1.01	3.73
89773			4.19	4.06	4.13	4.55	1.60	3.71
159401			3.80	3.42	3.34	3.87	4.10	3.71
50656	4		5.12	3.30	3.70	2.28	4.10	3.70
82491		1 1	4.19	3.64	3.11	2.99	4.52	3.69
8495			5.70	4.55	4.49	1.79	1.92	3.69
95108) i	6.73	4.60	2.27	2.30	2.53	3.69 3.68
3042	1		0.76	1.72	2.50	10.04	3.40	3.68
78409	E .		3.21	4.16	4.32	2.79	3.93	3.68
13590			9.57	1.87	4.30	1.31	1.34	3.68
95068	1		4.34	2.16	2.56	4.14	5.19	3.66
8782	2		5.36	2.63	2.49	1.62	6.22	3.66
84530	•		4.04	3.09	2.09	5.22	3.87	3.66
3584	•	1 1	4.06	4.52	2.15	4.94	2.64 2.69	3.65
762:			5.18	3.55	3.64	3.20	2.02	3.65
366	•		4.19	3.70	2.34	5.99	1.62	3.64
6242			3.97	2.76	2.08	7.77	1.87	3.63
7430	71 1.8	9 0.52	4.55	1.78	1.17	8.79	1.07	5.05

	1	ا مما		2.25	0.84	5.24	7.71	3.63
1070015	3.26	0.90	1.10	3.25 2.71	2.51	4.70	5.27	3.63
203132	12.63	3.48	2.95	3.86	2.56	4.66	3.46	3.63
259953	1.46	0.40	3.59	3.80 3.80	3.96	5.65	0.91	3.62
814427	0.47	0.13	3.78	1.43	1.22	13.31	1.06	3.62
259462	3.69	1.02	1.07	3.34	3.00	3.69	5.66	3.61
810558	14.49	4.01	2.38	3.3 4 4.60	2.81	5.13	2.49	3.61
51151	3.26	0.90	3.04	4.60 5.64	2.94	2.81	2.70	3.61
272750	20.57	5.70	3.98	3.64 2.71	5.71	1.52	3.03	3.61
360885	40.29	11.17	5.08	6.19	3.18	2.98	2.71	3.61
208082	7.90	2.19	2.97	1.36	6.42	4.29	4.83	3.61
627428	24.22	6.72	1.13		3.32	4.11	4.73	3.61
845345	5.09	1.41	2.54	3.32	2.51	5.34	4.83	3.60
417801	17.18	4.77	2.94	2.40 3.22	3.20	3.45	4.16	3.60
361922	9.18	2.55	3.94	3.22 3.99	3.53	4.38	3.77	3.60
898206	1.79	0.50	2.31	3.99 3.55	5.68	1.98	4.01	3.59
431296	1.85	0.51	2.75	2.83	3.47	7.37	3.44	3.59
587595	559.22	155.61	0.86	2.63 3.58	2.47	2.92	4.10	3.59
1632161	39.75	11.08	4.86	3.38 3.37	3.11	3.53	5.26	3.58
121196	22.80	6.36	2.64		2.21	3.08	3.60	3.58
823819	11.96	3.34	4.72	4.29	3.29	3.04	3.47	3.58
1456721	12.82	3.58	3.28	4.85	3.33	2.21	2.73	3.58
1500324	6.77	1.89	2.74	6.90	3.33 3.48	3.18	3.96	3.58
86160	2.10	0.59	3.16	4.11	3.40 3.47	5.00	1.89	3.58
246620	16.85	4.71	3.66	3.89	2.60	6.50	3.64	3.58
52730	6.16	1.72	2.03	3.11	2.52	2.52	4.76	3.58
809657	4.72	1.32	5.12	2.96	2.32	2.88	3.54	3.57
253577	1.48	0.41	4.99	4.45 3.80	3.21	5.49	2.40	3.57
486459	6.24	1.75	2.96	3.80 4.47	3.21	4.26	3.17	3.57
23576		1.00	2.58 3.95	4.47	2.67	3.43	3.62	3.57
119403	1.22	0.34	3.93 4.11	3.07	4.52	3.34	2.80	3.57
324342	1 (1.24	1.93	4.33	4.07	2.68	4.81	3.56
447518	1 1	3.08	1.93 3.87	4.33 4.29	4.07	2.34	2.86	3.55
811953		5.02	3.87 4.84	3.47	4.24	4.08	1.11	3.55
824124		0.29	4.84 2.00	4.04	3.97	5.08	2.63	3.55
837953	8	2.95	2.43	4.75	3.58	3.11	3.86	3.54
511521			2.43 3.88	4.73	1.98	5.20	2.16	3.54
1493252			2.29	3.09	5.39	3.53	3.42	3.54
626640	1		3.38	2.26	3.14	6.00	2.92	3.54
22731	1		5.71	3.40	2.86	3.42	2.30	3.54
84023	1		2.18	2.19	10.28	1.46		3.54
841303 150263		1 1	3.31	4.40	3.92	2.96	3.05	3.53
95046	· i		0.81	1.10	5.41	4.59	5.73	3.53
25101	1		2.89	2.88	6.72	1.14	4.00	3.52
25101 6765			1.22	3.32	2.33	6.57	4.11	3.51
			0.83	2.74	2.70	8.19	3.04	3.50
59240 75575	- 1	1	3.23	1.83	2.12	7.95	2.36	3.50
142079		1 1	2.46	3.91	3.17	3.47	4.47	3.49
84335	1	1 1	2.97	3.18	2.69	3.78	4.80	3.48
160434			1.56	2.15	1.34	9.00	3.33	3.48
25810		1 1	3.26	5.42	3.43	2.37	2.90	3.48
23010	11.2	1 4.50	5.20		2		. •	

308466	9.26	2.67	4.61	4.96	3.31	1.52	2.95	3.47
272576	6.22	1.80	1.34	3.85	4.11	4.71	3.25	3.45
108265	7.24	2.10	4.59	3.00	3.05	3.84	2.76	3.45
842784	326.54	94.88	0.85	3.11	1.95	7.01	4.30	3.44
1467207	3.68	1.07	2.33	4.23	4.08	2.71	3.86	3.44
784225	10.62	3.09	2.36	4.44	3.93	2.57	3.89	3.44
430179	51.95	15.13	2.52	2.04	9.04	1.80	1.78	3.44
796513	17.83	5.20	4.20	3.32	2.84	5.24	1.56	3.43
83363	5.08	1.48	3.97	3.58	1.64	3.63	4.33	3.43
489109	7.44	2.17	2.33	3.11	5.81	2.19	3.71	3.43
812977	7.62	2.22	2.38	2.15	6.48	4.79	1.35	3.43
1492238	13.96	4.08	4.19	3.72	2.61	4.20	2.41	3.43
740914	7.20	2.11	5.02	1.70	6.97	1.58	1.84	3.42
1475797	85.21	24.92	1.93	2.41	2.93	7.69	2.14	3.42
897042	4.95	1.45	6.12	3.62	2.70	3.57	1.07	3.42
249705	62.65	18.34	2.89	3.73	2.52	4.92	3.02	3.42
1476065		1.86	2.32	4.22	1.08	8.04	1.42	3.41
251731	0.82	0.24	2.88	3.28	3.37	3.75	3.78 4.57	3.41
8 38366	1 1	3.27	2.46	3.30	3.45	3.28		3.41 3.41
855799		2.64	2.70	3.06	2.67	2.66	5.96	3.41
701112	1 1	2.55	2.26	4.01	3.63	4.37	2.78	3.41 3.41
26406		1.92	2.41	3.58	3.68	3.80	3.58 2.30	3.41
727251		10.07	2.95	3.27	4.52	4.01	2.54	3.41
252453	1 1	5.91	2.32	4.20	2.91	5.06 3.49	2.08	3.40
1435185	1	0.75	5.55	2.80	3.10	5.49 5.40	3.13	3.40
741474	1	1 1	2.81	2.16	3.51 7.51	1.93	1.56	3.40
28989	3		3.60	2.40 2.86	3.04	3.54	4.02	3.40
79677			3.55	3.05	2.60	3.37	3.27	3.40
29064	1		4.70 2.27	1.81	4.99	1.02	6.89	3.40
14901	t .		7.43	1.72	1.39	4.51	1.94	3.40
74529	1	1 1	1.95	2.17	2.24	8.88	1.70	3.39
95060	1	1 1	3.07	4.92	1.88	5.02	2.03	3.38
87840 95043	li .		4.11	4.50	2.71	2.81	2.76	3.38
3982		1 1	2.98	3.80	3.61	3.36	3.14	3.38
84313			4.71	1.39	4.05	2.96	3.77	3.38
13718	1	1 1	2.28	4.02	3.12	4.62	2.80	3.37
29652	1	1 1	3.71	4.18	2.72	2.15	4.08	3.37
54463			4.23	3.65	3.42	1.95	3.59	3.37
84114	1	1 1	4.27	4.30	2.28	3.23	2.75	3.36
497			4.06	3.75	2.82	2.96	3.23	3.36
16388		1 1	3.84	4.80	3.21	2.46	2.52	3.36
510			3.25	1.40	2.40	2.89	6.87	3.36
7530		1 .	2.28	4.55	3.07	4.01	2.88	3.36
8982			2.34	3.18	3.05	4.51	3.70	3.36
4907	3	5 1.60	2.63	3.40	2.73	6.39	1.60	3.35
8556	1		1.02	1.19	0.83	1.13	12.53	3.34
7683	95 14.3	29 4.28	4.43	3.97	3.60	2.60	2.09	3.34
2121	65 41.5		5.13	3.22	4.38	1.81	2.15	3.34
3772	46 12.0	1 1	2.94	2.21	9.10	0.91	1.52	3.34
8098	376 27.	74 8.32	4.70	3.46	1.68	2.47	4.37	3.34

const	0.76	2 (2)	2.60	0.87	2.51	7.67	2.81	3.33
462325	8.76 6.17	2.63 1.86	2.80	3.81	2.96	4.60	1.86	3.32
624785		l l	3.38	1.97	3.76	1.35	1.64	3.32
291345	70.40	21.23	7.87		5.70 5.42	2.19	2.05	3.32
257382	12.80	3.86	3.91	3.02		3.74	3.40	3.31
700900	9.94	3.00	3.26	2.88	3.29 3.03	3.74	3.14	3.31
742581	1.20	0.36	3.04	4.31		4.66	2.46	3.31
811867	24.57	7.43	3.72	3.16	2.52			
502622	18.65	5.65	2.32	2.97	3.27	5.80	2.16	3.30 3.30
666371	0.85	0.26	3.76	2.72	3.71	3.35	2.97	
362829	3.03	0.92	3.18	5.12	3.27	3.18	1.73	3.30
129146	7.97	2.42	3.78	3.62	2.54	2.98	3.55	3.30
135083	17.07	5.19	3.09	3.47	4.93	2.81	2.15	3.29
854678	6.21	1.89	1.76	9.64	1.93	1.21	1.90	3.29
321470	7.74	2.36	3.41	4.28	3.25	1.89	3.60	3.29
1472638	7.11	2.16	2.45	3.79	2.73	4.52	2.93	3.29
242952	16.94	5.16	3.84	3.11	3.39	3.54	2.54	3.28
884657	6.68	2.04	3.32	2.87	2.28	4.95	2.99	3.28
40338	2.61	0.80	4.02	2.85	3.77	2.65	3.10	3.28
461670	8.90	2.71	2.03	4.16	4.85	2.37	2.99	3.28
45564	1.66	0.51	2.98	5.39	3.62	1.74	2.64	3.28
813637	34.08	10.41	3.65	3.22	2.20	5.54	1.75	3.27
291255	15.20	4.65	3.07	3.01	5.02	2.76	2.48	3.27
310493	7.56	2.32	3.61	3.65	3.06	2.77	3.24	3.26
433350	28.82	8.83	1.93	2.66	2.67	7.86	1.21	3.26
781454	9.75	2.99	2.79	1.30	9.94	1.22	1.06	3.26
685516	3.36	1.03	3.63	3.79	2.76	4.08	2.03	3.26
112785	7.75	2.38	2.21	3.83	3.58	3.82	2.81	3.25
264640	13.22	4.07	3.98	3.82	1.83	1.43	5.20	3.25
202904	12.77	3.93	2.44	4.34	2.65	3.34	3.48	3.25
251330	9.21	2.84	2.30	2.37	2.73	4.39	4.44	3.25
815164	0.88	0.27	3.76	3.41	4.07	3.68	1.32	3.25
1456602	6.11	1.88	2.40	2.64	2.44	4.72	4.03	3.25
1486260	13.23	4.08	2.75	2.87	3.17	5.07	2.36	3.24
322561	107.98	33.31	2.93	2.03	1.79	4.79	4.67	3.24
416436	11.08	3.42	5.36	3.7 3	2.74	3.15	1.23	3.24
827152	3.50	1.08	1.39	5.25	3.59	4.05	1.92	3.24
289818	7.47	2.31	2.49	4.60	3.59	3.30	2.19	3.23
273546	9.10	2.81	4.41	3.56	2.54	2.67	2.98	3.23
75886	14.08	4.37	5.08	3.68	1.09	4.18	2.11	3.23
416010	53.71	16.66	2.41	1.81	8.72	1.50	1.69	3.23
838628	2.82	0.88	2.12	3.44	3.70	3.20	3.67	3.23
2274	7 37.20	11.55	2.96	3.25	2.59	4.14	3.15	3.22
95109	1 2.03	0.63	2.71	3.17	4.10	2.80	3.30	3.22
3581	2 2.29	0.71	2.94	2.49	2.99	3.34	4.32	3.22
8287	4 5.14	1.60	4.36	3.56	2.21	3.84	2.10	3.21
163171	3 5.68	1.77	2.81	3.64	3.90	2.77	2.93	3.21
26307	0 1.17	0.37	4.55	4.07	1.99	3.33	2.10	3.21
103169	8 6.26	1.95	2.99	2.12	2.54	4.81	3.58	3.21
160212	0 6.44	2.01	1.98	3.03	2.64	5.10	3.30	3.21
45467	7.6	8 2.40	5.39	3.57	1.75	2.15	3.16	3.20
83790	103.1	32.18	2.10	2.12	2.01	4.30	5.50	3.20

34869	2.74	0.85	1.85	2.96	1.76	7.56	1.87	3.20
840590	21.70	6.78	2.44	1.18	2.39	8.32	1.68	3.20
377672	2.59	0.81	0.93	0.98	0.68	12.46	0.96	3.20
511632	7.85	2.46	2.42	2.96	2.15	4.52	3.94	3.20
531028	21.14	6.62	3.62	3.65	3.05	2.41	3.26	3.20
321386	2.45	0.77	3.26	2.66	2.50	4.45	3.11	3.20
754221	6.80	2.13	2.49	3.95	5.19	2.42	1.93	3.20
433573	10.67	3.34	3.00	3.46	3.80	2.33	3.38	3.19
489823	33.49	10.49	2.75	4.23	1.58	5.36	2.05	3.19
248454	3.70	1.16	7.24	2.69	2.10	2.69	1.24	3.19
298155	2.85	0.89	2.52	4.31	2.59	3.88	2.66	3.19
530814	46.55	14.60	0.83	4.74	4.23	4.56	1.60	3.19
294259	4.21	1.32	3.87	2.34	3.12	1.09	5.52	3.19
726488	10.10	3.17	3.31	3.38	3.27	3.50	2.45	3.18
511459	37.04	11.64	2.30	3.42	4.76	3.38	2.06	3.18
503468	51.40	16.15	3.74	2.80	1.75	4.34	3.28	3.18
756600	47.06	14.79	3.37	1.74	1.79	4.21	4.80	3.18
760344	5.89	1.85	3.62	4.43	2.41	1.42	4.03	3.18
753252	7.78	2.45	2.52	2.78	2.96	4.05	3.59	3.18
26230	5.08	1.60	1.18	3.71	4.40	3.50	3.09	3.18
37980	0.84	0.27	1.41	2.73	1.73	8.06	1.95	3.18
156386	2.14	0.67	3.02	3.99	3.91	2.87	2.09	3.18
969844	9.12	2.88	5.15	2.28	2.19	3.92	2.33	3.17
855723	6.73	2.12	2.86	3.79	2.80	3.00	3.38	3.17
1240220	7.60	2.40	0.79	2.71	6.02	4.99	1.31	3.17
451769	5.25	1.66	3.67	2.29	2.52 .	5.57	1.78	3.16
810027	0.44	0.14	3.75	3.05	2.90	3.82	2.31	3.16
731272		1.59	2.45	2.89	3.95	4.89	1.62	3.16
129024		1.36	5.53	0.58	1.77	1.86	6.04	3.16 3.15
795907		1.71	1.43	1.64	3.37	0.61	8.72 1.45	3.15
731257	1 1	1.41	1.78	2.17	3.22	7.14	3.63	3.15
703707		1.31	2.68	2.65	3.51	3.30 3.41	3.03	3.15
811603		6.22	3.87	2.99	2,25 3.45	3.41	2.15	3.15
884346		1.71	2.20	4.78 2.46	3.43 1.74	5.00	2.35	3.14
201628		2.39 5.60	4.15	3.10	3.72	2.88	2.71	3.14
840821	1 1	1	3.31 3.94	2.68	2.53	2.44	4.11	3.14
344432 590853	1 1	0.21	2.61	2.78	3.43	3.40	3.47	3.14
767298	1 1	6.18	2.43	3.33	2.32	4.12	3.48	3.14
3638	1 1		2.66	3.88	4.50	1.19	3.46	3.14
30295		3.39	1.90	5.11	2.20	4.12	2.36	3.14
79688			4.15	1.95	2.26	4.54	2.77	3.13
72548		8.54	3.93	3.59	2.81	3.60	1.72	3.13
12148	1 !		3.79	3.88	3.08	2.54	2.35	3.13
26389	1		3.70	3.08	2.13	4.71	2.03	3.13
78238	1		2.40	2.83	2.35	5.64	2.42	3.13
81556	4		3.37	3.78	2.43	4.15	1.90	3.13
24554	E .	1	1.15	5.91	3.40	3.50	1.66	3.12
85657	1	1 1	2.42	1.67	5.16	3.07	3.31	3.12
25175	t t		2.40	2.73	2.75	5.27	2.45	3.12
35699		7.04	4.19	2.74	2.51	3.63	2.54	3.12
	-	•						

502463	5.77	1.85	2.78	3.60	3.10	2.63	3.49	3.12
1461528	3.95	1.27	3.05	2.21	5.36	2.64	2.33	3.12
244652	24.70	7.92	2.62	1.81	7.76	1.54	1.87	3.12
266106	20.68	6.63	2.76	3.21	6.30	1.40	1.93	3.12
1415981	4.13	1.32	3.58	3.02	4.59	2.33	2.07	3.12
66902	3.21	1.03	4.24	2.05	1.56	0.87	6.87	3.12
283919	7.56	2.43	7.31	3.63	1.47	2.02	1.15	3.11
813712	7.62	2.45	1.14	3.64	1.91	6.70	2.18	3.11
195903	6.95	2.24	2.45	4.56	3.51	2.97	2.05	3.11
345032	20.78	6.69	1.19	4.73	3.52	2.65	3.46	3.11
1635665	5.13	1.65	4.57	3.12	0.90	5.26	1.69	3.11
154749	1.76	0.57	3.34	3.27	3.77	2.57	2.58	3.11
287728	2.76	0.89	0.59	4.01	0.98	9.25	0.70	3.11
701409	1.54	0.50	4.21	2.63	3.95	2.39	2.34	3.10
627687	56.19	18.11	1.06	1.86	3.60	4.82	4.16	3.10
855244	15.78	5.09	6.04	0.82	7.22	0.56	0.88	3.10
293243	13.17	4.25	4.44	2.03	6.12	0.95	1.96	3.10
295206	3.16	1.02	5.72	3.11	1.43	2.48	2.76	3.10
744846	2.69	0.87	3.69	4.51	2.68	3.60	1.01	3.10
251645	0.50	0.16	3.12	3.21	3.74	3.30	2.12	3.10
357364	0.72	0.23	5.76	3.07	1.81	3.15	1.68	3.09
451707	7.59	2.46	2.46	1.83	6.35	1.34	3.48	3.09
250673	2.66	0.86	7.23	1.34	2.15	1.85	2.88	3.09
1031086	25.15	8.15	3.21	2.15	2.25	5.00	2.83	3.09
50892	7.95	2.58	3.62	2.53	3.85	2.69	2.71	3.08
52327	5.70	1.85	1.94	3.44	2.70	3.46	3.87	3.08
448930	7.53	2.45	2.47	2.25	2.41	3.62	4.66	3.08
713109	2.44	0.79	1.97	2.95	3.60	4.59	2.28	3.08
815794	14.69	4.77	5.65	1.93	1.67	1.95	4.18	3.08
789091	1 1	3.09	6.20	4.45	1.91	1.88	0.95	3.08
950836	1 1	11.31	2.78	1.53	8.20	0.92	1.95	3.08
213890		4.60	2.37	5.07	2.41	3.28	2.24	3.07
74223		1.76	3.29	3.47	3.41	2.68	2.52	3.07
41174		1.31	3.25	3.60	2.82	3.70	1.98	3.07
1635874		1.97	6.28	3.19	1.83	2.38	1.69	3.07
810284		0.74	5.17	2.15	2.18	2.13	3.71	3.07
845502	1 1	1.26	1.42	3.27	3.03	4.57	3.06	3.07 3.07
461489		0.30	3.19	3.40	2.48	2.98	3.28 3.64	3.07
36109	, ,	1.84	3.06	3.81	1.75	3.06	1.10	3.06
13911	1 1	1.41	2.39	3.80	2.88	5.14 4.58	2.11	3.06
81363		27.59	3.51	2.53	2.58 2.21	4.36 3.39	3.38	3.06
74380		1.66	3.74	2.59	2.21	2.50	3.06	3.06
48656		2.99	2.48	4.29	2.98	2.65	2.73	3.06
66512			3.90	3.70	2.86	2.82	5.58	3.06
13660		1 1	3.42 2.38	0.62 3.22	3.22	3.25	3.22	3.06
39562		3 t	2.38 3.56	3.22 0.96	5.02	3.25 3.66	2.06	3.06
28752 32347	1		5.34	3.25	2.05	2.36	2.27	3.06
3234 / 88438	1	5 t	2.97	2.78	2.75	3.38	3.40	3.06
22252	1		1.68	4.03	3.69	4.09	1.79	3.05
26062	l l	1 1	1.68	4.70	3.88	2.83	2.17	3.05
2000	,	. ~., ~!			5.03			

471266 2.72 0.89 3.15 2.95 2.66 3.45 3.03 3.05 565062 1.38 0.45 1.72 3.15 3.97 2.87 3.53 3.05 878127 3.53 1.16 2.07 3.35 4.53 3.32 1.96 884766 9.95 3.13 1.03 2.09 3.44 2.70 5.00 2.00 3.05 884766 9.95 3.27 1.22 3.47 2.45 2.67 5.40 3.04 754192 3.0 1.09 4.21 2.70 2.65 3.45 2.21 3.04 344589 53.28 17.55 1.28 4.11 3.84 2.63 3.31 3.04 328713 0.46 0.15 2.60 3.52 3.31 2.33 3.42 3.03 328718 1.57 3.43 1.95 3.39 2.83 3.03 328718 3.44 1.79 3.47 2.18<									
701532	471266	2.72	0.89	3.15	2.95	2.66	3.45	3.03	3.05
878127 3.53 1.16 2.07 3.35 4.53 3.32 1.96 3.05 878373 31.06 10.19 3.07 3.84 2.03 4.00 2.29 3.05 884766 9.95 3.27 1.122 3.47 2.45 2.67 5.40 3.04 754192 3.30 1.09 4.21 2.70 2.65 3.45 2.21 3.04 344589 53.28 17.55 1.28 4.11 3.84 2.63 3.31 3.04 344589 53.28 17.55 3.27 3.89 4.14 2.12 1.76 3.04 328613 0.46 0.15 2.60 3.52 3.31 2.33 3.42 3.03 388467 24.71 8.14 3.57 3.43 1.95 3.39 2.83 3.03 828928 5.44 1.79 3.47 2.18 3.94 3.62 3.03 3.03 3.03 3.03 3.03 <	565062	1.38	0.45	1.72	3.15	3.97	2.87	3.53	3.05
878373 31.06 10.19 3.07 3.84 2.03 4.00 2.29 3.05 40075 3.13 1.03 2.09 3.44 2.70 5.00 2.00 3.05 844766 9.95 3.27 1.22 3.47 2.45 2.67 5.40 3.04 754192 3.30 1.09 4.21 2.70 2.65 3.45 2.21 3.04 39154 1.37 0.45 1.64 3.95 3.01 2.17 4.42 3.04 487153 10.77 3.55 3.27 3.89 4.14 2.12 1.76 3.04 887153 10.77 3.55 3.27 3.89 4.14 2.12 1.76 3.03 3280288 5.44 1.79 3.47 2.18 3.94 1.96 3.62 3.03 382818 13.97 4.61 2.21 4.58 2.41 3.19 2.78 3.03 383130 11.12 3	701532	3.26	1.07	3.06	2.86	3.51	1.98	3.84	3.05
40075 3.13 1.03 2.09 3.44 2.70 5.00 2.00 3.05 884766 9.95 3.27 1.122 3.47 2.45 2.67 5.40 3.04 754192 3.30 1.09 4.21 2.70 2.65 3.45 2.21 3.04 299154 1.37 0.45 1.64 3.95 3.01 2.17 4.42 3.04 3444589 53.28 17.55 1.28 4.11 3.84 2.63 3.31 3.04 328613 0.46 0.15 2.60 3.52 3.31 2.33 3.42 3.03 884867 24.71 8.14 3.57 3.43 1.95 3.39 2.83 3.03 828842 6.00 1.98 1.78 2.66 4.19 1.45 5.08 3.03 825842 6.00 1.98 1.78 2.66 4.19 1.45 5.08 3.03 133130 1.12 3.	878127	3.53	1.16	2.07	3.35	4.53			
884766 9.95 3.27 1.22 3.47 2.45 2.67 5.40 3.04 754192 3.30 1.09 4.21 2.70 2.65 3.45 2.21 3.04 299154 1.37 0.45 1.64 3.95 3.01 2.17 4.42 3.04 344589 53.28 17.55 1.28 4.11 3.84 2.63 3.31 3.04 897153 10.77 3.55 3.27 3.89 4.14 2.12 1.76 3.04 328613 0.46 0.15 2.60 3.52 3.31 2.33 3.03 3.03 884867 24.71 8.14 3.57 3.43 1.95 3.39 2.83 3.03 8825842 6.00 1.98 1.78 2.66 4.19 1.45 5.08 3.03 825842 6.00 1.98 1.78 2.66 4.19 1.45 5.08 3.03 825842 1.00 0	878373	31.06	10.19	3.07	3.84	2.03	4.00		
754192 3.30 1.09 4.21 2.70 2.65 3.45 2.21 3.04 299154 1.37 0.45 1.64 3.95 3.01 2.17 4.42 3.04 34589 53.28 17.55 12.8 4.11 3.84 2.63 3.31 3.04 397133 10.77 3.55 3.27 3.89 4.14 2.12 1.76 3.04 328613 0.46 0.15 2.60 3.52 3.31 2.33 3.42 3.03 884867 24.71 8.14 3.57 3.43 1.95 3.39 2.83 3.03 2.85 3.42 1.79 3.47 2.18 3.94 1.96 3.62 3.03 3.85 28528 5.44 1.79 3.47 2.18 3.94 1.96 3.62 3.03 328528 5.44 1.79 3.47 2.18 3.94 1.96 3.62 3.03 313103 11.12 3.67 3.58 0.72 2.94 2.70 5.20 3.03 313130 11.12 3.67 3.58 0.72 2.94 2.70 5.20 3.03 813187 22.70 7.51 1.31 2.52 4.98 3.97 2.34 3.02 2.6564 1.20 0.40 2.46 3.75 2.39 3.50 3.03 3.03 813187 22.70 7.51 1.31 2.52 4.98 3.97 2.34 4.50 2.40 3.02 2.46 2.73 3.02 2.66747 3.05 1.01 3.26 3.57 2.70 0.99 4.59 3.02 2.66747 3.05 1.01 3.26 3.57 2.70 0.99 4.59 3.02 2.52845 9.34 3.10 4.67 3.03 3.85 2.20 2.08 4.19 2.75 3.02 3.22914 9.34 3.10 4.67 3.02 2.07 1.93 3.39 3.02 445009 15.29 5.07 3.85 3.88 2.20 2.08 4.19 2.75 3.02 3.22914 9.34 3.10 4.67 3.02 2.07 1.93 3.39 3.02 4.6708 4.29 1.42 1.21 3.61 4.50 3.23 2.53 3.02 4.50 3.03 3.03 3.03 3.03 3.03 3.03 3.03 3	40075	3.13	1.03	2.09	3.44	2.70			
299154	884766	9.95	3.27	1.22	3.47	2.45		•	
344589 53.28 17.55 1.28 4.11 3.84 2.63 3.31 3.04 897153 10.77 3.55 3.27 3.89 4.14 2.12 1.76 3.04 328613 0.46 0.15 2.60 3.52 3.31 2.33 3.42 384867 24.71 8.14 3.57 3.43 1.95 3.39 2.83 3.03 3825842 6.00 1.98 1.78 2.66 4.19 1.45 5.08 3.03 3525842 6.00 1.98 1.78 2.66 4.19 1.45 5.08 3.03 3528644 1.20 0.40 2.46 3.75 2.39 3.50 3.03 3513187 22.70 7.51 1.31 2.52 4.98 3.97 2.34 3.02 252864 1.20 0.40 2.46 3.75 2.39 3.50 3.03 3.03 313183187 2.20 0.40 2.46	754192	3.30	1.09	4.21	2.70	2.65			
897153 10.77 3.55 3.27 3.89 4.14 2.12 1.76 3.04 328613 0.46 0.15 2.60 3.52 3.31 2.33 3.42 3.03 884867 24.71 8.14 3.57 3.43 1.95 3.39 2.83 3.03 885818 13.97 4.61 2.21 4.58 2.41 3.19 2.78 3.03 825842 6.00 1.98 1.78 2.66 4.19 1.45 5.08 3.03 313130 11.12 3.67 3.58 0.72 2.94 2.70 5.20 3.03 313187 22.70 7.51 1.31 2.52 4.98 3.97 2.34 3.02 2.46 2.73 3.02 2.66747 3.05 1.01 3.26 3.57 2.70 0.99 4.59 3.02 460747 3.05 1.01 3.26 3.57 2.70 0.99 4.59 3.02 <td< td=""><td>299154</td><td>1.37</td><td>0.45</td><td>1.64</td><td>3.95</td><td></td><td></td><td></td><td></td></td<>	299154	1.37	0.45	1.64	3.95				
328613 0.46 0.15 2.60 3.52 3.31 2.33 3.42 3.03 884867 24.71 8.14 3.57 3.43 1.95 3.39 2.83 3.03 289288 5.44 1.79 3.47 2.18 3.94 1.96 3.62 3.03 825842 6.00 1.98 1.78 2.66 4.19 1.45 5.08 3.03 825842 6.00 1.98 1.78 2.66 4.19 1.45 5.08 3.03 825864 1.20 0.40 2.46 3.75 2.39 3.50 3.03 3.03 813187 22.70 7.51 1.31 2.52 4.98 3.97 2.34 3.02 1635110 38.13 1.262 4.50 2.40 3.02 2.46 2.73 3.02 4690360 5.94 1.97 2.87 3.33 2.69 3.50 2.72 3.02 951216 2.02	344589	53.28	17.55	1.28	4.11	3.84		1	
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360232 2.20 0.74 1.98 3.91 3.63 3.66 1.72 2.98 74511 5.23 1.75 4.10 3.44 2.52 0.97 3.89 2.98 343167 15.62 5.24 3.15 4.46 2.96 2.05 2.27 2.98 39885 0.94 0.31 0.88 2.80 4.66 4.76 1.80 2.98 122183 7.61 2.56 2.66 4.22 2.01 2.53 3.46 2.98 38253 2.80 0.94 3.19 4.23 2.69 2.79 1.98 2.98 745103 8.87 2.98 2.03 3.44 3.39 2.05 3.96 2.97 155838 2.53 0.85 2.70 1.94 5.31 2.18 2.74 2.97									
74511 5.23 1.75 4.10 3.44 2.52 0.97 3.89 2.98 343167 15.62 5.24 3.15 4.46 2.96 2.05 2.27 2.98 39885 0.94 0.31 0.88 2.80 4.66 4.76 1.80 2.98 122183 7.61 2.56 2.66 4.22 2.01 2.53 3.46 2.98 38253 2.80 0.94 3.19 4.23 2.69 2.79 1.98 2.98 745103 8.87 2.98 2.03 3.44 3.39 2.05 3.96 2.97 155838 2.53 0.85 2.70 1.94 5.31 2.18 2.74 2.97			1					T .	
343167 15.62 5.24 3.15 4.46 2.96 2.05 2.27 2.98 39885 0.94 0.31 0.88 2.80 4.66 4.76 1.80 2.98 122183 7.61 2.56 2.66 4.22 2.01 2.53 3.46 2.98 38253 2.80 0.94 3.19 4.23 2.69 2.79 1.98 2.98 745103 8.87 2.98 2.03 3.44 3.39 2.05 3.96 2.97 155838 2.53 0.85 2.70 1.94 5.31 2.18 2.74 2.97		3							
39885 0.94 0.31 0.88 2.80 4.66 4.76 1.80 2.98 122183 7.61 2.56 2.66 4.22 2.01 2.53 3.46 2.98 38253 2.80 0.94 3.19 4.23 2.69 2.79 1.98 2.98 745103 8.87 2.98 2.03 3.44 3.39 2.05 3.96 2.97 155838 2.53 0.85 2.70 1.94 5.31 2.18 2.74 2.97		1	4						
122183 7.61 2.56 2.66 4.22 2.01 2.53 3.46 2.98 38253 2.80 0.94 3.19 4.23 2.69 2.79 1.98 2.98 745103 8.87 2.98 2.03 3.44 3.39 2.05 3.96 2.97 155838 2.53 0.85 2.70 1.94 5.31 2.18 2.74 2.97									
38253 2.80 0.94 3.19 4.23 2.69 2.79 1.98 2.98 745103 8.87 2.98 2.03 3.44 3.39 2.05 3.96 2.97 155838 2.53 0.85 2.70 1.94 5.31 2.18 2.74 2.97			1						
745103 8.87 2.98 2.03 3.44 3.39 2.05 3.96 2.97 155838 2.53 0.85 2.70 1.94 5.31 2.18 2.74 2.97			1 1						
155838 2.53 0.85 2.70 1.94 5.31 2.18 2.74 2.97		1	1 1						
		L	1 1						
813143 0.11 2.00 3.31 3.70 2.01 2.36 1.21 2.37									
	81314	ا۱.0 الا	11 2.00	3.37	3.70	2.01	2.30	1.21	2.71

469685	4.95	1.67	3.25	3.34	2.23	3.52	2.51	2.97
824640	6.96	2.35	1.37	1.54	3.69	3.65	4.59	2.97
825833	1.95	0.66	3.29	2.75	4.19	2.65	1.97	2.97
25517	8.52	2.88	3.15	3.29	2.67	2.43	3.28	2.97
842767	2.15	0.73	2.59	3.85	3.19	2.67	2.53	2.96
148225	3.89	1.31	2.29	2.66	1.88	5.50	2.47	2.96
772304	113.14	38.24	3.78	2.19	2.18	2.83	3.83	2.96
246800	6.67	2.26	2.06	3.22	2.26	4.33	2.91	2.96
322192	7.97	2.69	1.72	3.28	5.02	1.74	3.02	2.96
755765	4.01	1.36	2.43	3.91	4.37	1.46	2.62	2.96
462468	3.24	1.10	3.23	3.74	3.40	1.72	2.69	2.96
701402	4.23	1.43	1.82	3.11	5.63	2.91	1.31	2.96
253253	1.73	0.59	4.71	4.49	1.70	2.17	1.68	2.95
211206	4.71	1.60	2.58	3.21	2.52	2.63	3.82	2.95
813825	0.52	0.18	3.85	3.04	1.69	4.84	1.32	2.95
898227	2.02	0.69	1.45	4.11	2.91	4.09	2.18	2.95
45578	5.05	1.72	4.61	2.16	2.26	3.42	2.27	2.94
1522799		0.42	2.25	3.40	2.72	3.47	2.86	2.94
593520		2.13	2.05	2.67	3.01	4.62	2.34	2.94
487773		10.06	4.78	2.56	4.05	1.34	1.95	2.93
502690	1 1	1.39	2.22	3.28	2.51	4.37	2.29	2.93
731104		0.22	3.42	2.35	3.12	2.70	3.07	2.93
796341		2.44	2.21	2.93	2.48	3.55	3.49	2.93
1492304	3	1.44	2.24	2.81	3.19	4.05	2.36	2.93
77805	1 1	5.31	2.32	4.51	2.41	2.27	3.13	2.93
415700		1.15	2.29	2.89	4.79	1.53	3.14	2.93
701496		0.78	2.43	2.22	5.16	3.20	1.62	2.93
233419		1.90	1.88	3.96	3.98	1.76	3.06	2.93 2.93
767994		1.47	2.76	1.52	8.75	0.69	0.92 1.00	2.93
254310		3.51	3.10	2.50	6.68	1.35 2.44	2.76	2.92
4803:		0.18	2.77	3.62 3.26	3.02 2.72	2. 44 2.81	3.05	2.92
81184		3.39	2.78	4.15	2.72	2.69	3.08	2.92
4497		3.05	2.47 2.27	2.18	2.15	5.77	2.23	2.92
24306		1	2.60	2.18	5.83	2.20	1.33	2.92
163535	1	1	2.12	1.92	1.94	7.00	1.59	2.91
75994 3580			3.30	2.96	2.26	3.58	2.45	2.91
4551			1.58	3.77	5.25	1.91	2.04	2.91
159162	5		2.27	3.52	1.93	4.64	2.17	2.91
46084	1		4.19	1.78	4.04	2.42	2.11	2.91
141172	1		3.69	2.02	2.19	4.86	1.78	2.91
16048			3.02	1.25	1.91	2.64	5.72	2.91
74269		1 1	2.56	3.08	2.90	4.14	1.83	2.90
2427			3.05	3.96	1.69	3.81	1.99	2.90
8688			1.07	1.24	0.96	9.00	2.21	2.90
14208			1.37	2.47	2.52	4.60	3.53	2.90
8257			2.16	5.30	2.82	2.12	2.09	2.90
8261	1		2.02	1.86	7.90	1.13	1.56	2.89
4614		3 8	3.29	3.33	2.51	2.75	2.59	2.89
494	1		3.81	2.49	2.73	3.24	2.15	2.89
1297	25 17.9	6.22	2.93	1.99	4.17	3.16	2.17	2.89

627273	1.58	0.55	1.86	3.85	2.31	3.67	2.73	2.88
143306	10.42	3.62	3.88	2.52	1.98	3.46	2.55	2.88
730313	3.95	1.37	2.62	2.27	3.50	2.08	3.93	2.88
825718	4.03	1.40	2.44	2.50	3.77	3.02	2.67	2.88
1591264	3.18	1.11	4.68	4.17	2.19	2.11	1.23	2.88
784016	33.80	11.75	2.08	4.19	3.69	2.77	1.66	2.88
41940	10.51	3.65	1.79	2.81	3.34	3.64	2.80	2.88
841501	5.94	2.07	3.74	4.22	2.48	2.62	1.31	2.88
853968	12.10	4.21	3.42	3.85	5.08	1.19	0.84	2.87
489663	2.11	0.73	3.88	2.51	1.67	3.15	3.17	2.87
669371	4.99	1.74	2.07	1.98	7.30	1.15	1.87	2.87
815760	2.31	0.81	3.18	2.00	4.13	3.96	1.09	2.87
125722	3.28	1.14	4.52	2.27	2.75	3.95	0.86	2.87
897625	2.58	0.90	1.23	2.12	4.08	3.14	3.79	2.87
768179	3.80	1.32	1.92	3.06	2.43	4.11	2.83	2.87
768469	4.17	1.45	2.78	1.73	2.41	3.64	3.79	2.87
898265	20.16	7.03	1.31	2.31	3.96	2.43	4.34	2.87
868119	10.00	3.49	2.81	3.31	3.66	1.82	2.73	2.87
897983	26.99	9.42	2.98	3.14	2.24	2.33	3.64	2.86
1416086	8.24	2.88	3.59	2.09	5.52	1.57	1.54	2.86
141972	18.88	6.60	3.58	3.00	3.07	1.99	2.68	2.86
144977	71.81	25.09	2.67	3.73	3.36	1.42	3.14	2.86
949940	12.52	4.38	4.69	2.75	1.41	3.20	2.25	2.86
898251	2.02	0.71	2.79	2.93	3.12	4.01	1.45	2.86
80500	140.81	49.33	0.91	2.65	1.64	5.24	3.85	2.86
823588	7.72	2.70	2.44	3.65	3.22	2.51	2.45	2.85
269300	10.64	3.73	1.47	2.55	4.46	4.07	1.71	2.85
823627	3.25	1.14	2.72	3.08	3.03	1.90	3.53	2.85
1589973		1.81	2.56	3.21	1.99	4.51	1.99	2.85
246541	10.77	3.78	2.76	3.03	2.41	3.92	2.14	2.85
784253	1 1	4.12	2.36	3.67	2.76	2.44	3.04	2.85 2.85
27072		0.10	2.61	2.23	3.44	2.93	3.04	2.85 2.85
754449		0.63	2.10	3.63	2.52	4.00	1.99	2.85 2.85
82905		0.43	3.27	1.88	1.90	3.49	3.70	2.85
743828		1.17	1.16	1.69	5.22	0.99	5.18 1.72	2.84
743851		2.89	3.40	2.79	2.54	3.77 3.86	2.70	2.84
1475746		4.39	3.24	2.13	2.29	3.86 6.19	2.78	2.84
283196		1.34	1.51	2.93	1.19 2.96	2.62	3.33	2.84
866276		0.68	1.99	3.31 2.02	2.96 2.76	3.87	1.75	2.84
144926	1 1	3.24	3.79 6.19	1.95	2.76	2.39	1.09	2.83
769926		3.51 4.20	2.36	3.04	1.33	4.78	2.64	2.83
730554			2.97	3.32	1.84	3.86	2.12	2.82
4001° 89757°		0.18	2.18	2.74	3.06	3.81	2.31	2.82
149241			1.73	2.18	3.44	2.84	3.92	2.82
133626	1		4.41	2.34	4.62	1.32	1.41	2.82
25067	1		3.14	3.84	3.52	1.72	1.86	2.82
81022	1	1 1	4.33	3.00	1.34	3.50	1.92	2.82
27107	•	1 1	2.34	3.42	3.53	2.75	2.04	2.82
11181		1 1	3.30	4.08	4.25	1.15	1.29	2.81
89809	1	1 5	3.41	2.02	3.72	1.08	3.83	2.81

250869	123.92	44.10	8.11	2.05	1.68	0.76	1.46	2.81
840620	13.81	4.92	1.46	1.93	2.95	3.21	4.47	2.81
53165	0.89	0.32	3.53	3.16	2.83	2.33	2.16	2.80
784128	1.83	0.65	1.61	1.75	6.85	1.95	1.84	2.80
452668	4.95	1.77	2.13	1.80	2.80	5.84	1.44	2.80
811911	11.89	4.25	3.15	1.99	3.02	2.42	3.41	2.80
854122	4.20	1.50	5.92	3.38	2.72	0.79	1.18	2.80
491184	6.35	2.27	3.28	1.15	8.07	0.69	0.80	2.80
825356	1.18	0.42	2.48	2.51	4.19	3.57	1.22	2.79
214884	3.55	1.27	4.42	2.54	1.68	3.83	1.50	2.79
450711	1.29	0.46	2.21	3.64	2.22	4.08	1.82	2.79
1593502	2.25	0.81	1.62	3.22	2.57	2.01	4.54	2.79
825058	3.42	1.23	2.71	1.17	8.02	1.37	0.68	2.79
285226	25.74	9.23	7.63	1.20	2.23	1.77	1.12	2.79
191904	7.67	2.75	3.12	2.55	2.70	2.27	3.28	2.79
786609	4.81	1.73	0.70	0.71	2.56	9.16	0.80	2.78
204146	2.40	0.86	1.54	5.79	2.84	1.41	2.34	2.78
713286	0.89	0.32	3.32	2.65	3.34	3.52	1.08	2.78
416833	6.07	2.18	1.35	3.81	1.74	4.90	2.09	2.78
268211	3.43	1.23	2.33	1.79	1.77	1.26	6.74	2.78
782439	35.89	12.92	3.09	2.38	1.70	3.98	2.74	2.78
810741	72.43	26.10	3.64	2.64	2.20	3.00	2.38	2.78
431286	13.81	4.98	2.22	2.98	3.24	1.94	3.50	2.77
826335	2.24	0.81	2.09	2.46	3.66	3.16	2.49	2.77
796313		0.28	3.22	2.69	2.55	2.13	3.26	2.77
812989		0.79	2.58	2.17	3.23	3.95	1.93	2.77
258521	1	1.20	1.48	7.97	1.12	1.13	2.14	2.77
526945		0.41	3.17	1.74	2.47	3.11	3.36	2.77
197821	1 1	2.15	5.83	4.39	1.04	1.29	1.28	2.77
1631747		1.84	4.14	4.87	1.99	1.76	1.08	2.77
1475738	: .	86.12	3.33	2.08	2.23	3.93	2.25	2.76
248261		0.55	4.72	1.35	0.99	1.14	5.61	2.76
841348		4.33	4.50	2.03	0.87	4.69	1.72	2.76
1031346		1.16	3.61	2.85	2.26	3.16	1.91	2.76 2.76
244154		1.79	2.49	2.18	3.84	2.06	3.22 1.09	2.76 2.76
245860		11.02	5.36	1.24	5.40	0.69	3.07	2.76 2.76
271472		4.28 2.36	2.07	3.57	2.66 1.98	2.40 3.47	1.95	2.75
75391		1.33	2.95 2.28	3.41 1.01	1.83	2.34	6.28	2.75
5174	1 1	2.18	2.28 1.94	3.44	2.46	2.34 2.47	3.43	2.75
29642 ⁶ 43031			1.77	2.79	3.60	1.97	3.59	2.74
28314			3.20	3.03	3.74	1.95	1.78	2.74
43534		1 I	3.20	2.13	2.58	3.61	2.19	2.74
27853		1	1.66	3.41	1.66	4.44	2.52	2.74
48972	E .		2.33	2.25	3.98	3.06	2.07	2.74
81494	1		2.62	3.11	2.36	3.47	2.14	2.74
50374			1.17	3.06	1.97	5.66	1.82	2.74
163214	3		2.34	3.20	3.06	2.96	2.13	2.74
42821	B	1 1	3.87	3.41	1.92	2.28	2.19	2.74
8783			2.19	2.60	1.52	4.16	3.20	2.73
15730		1 1	1.63	2.09	4.06	4.16	1.72	2.73
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841399	9.08	3.33	2.44	2.07	3.48	3.21	2.45	2.73
897730	41.12	15.07	3.07	1.82	2.32	2.54	3.88	2.73
866882	10.73	3.94	1.81	4.01	3.71	1.83	2.28	2.73
1160558	8.03	2.94	2.15	3.29	2.53	3.66	2.00	2.73
127928	3.67	1.35	1.79	3.28	2.89	2.15	3.51	2.73
897971	11.55	4.24	1.43	2.82	3.44	2.74	3.21	2.73
447568	9.00	3.31	3.49	1.70	1.86	4.48	2.08	2.72
811880	3.13	1.15	1.85	2.58	4.75	3.12	1.30	2.72
810725	3.27	1.20	1.92	2.23	1.63	5.97	1.85	2.72
197176	20.79	7.65	3.06	2.51	3.48	2.16	2.38	2.72
758347	2.75	1.01	1.61	3.59	3.19	2.80	2.41	2.72
43705	0.35	0.13	2.72	3.74	1.90	3.12	2.09	2.71
813280	4.14	1.53	1.72	2.58	2.29	3.06	3.91	2.71
195652	18.03	6.65	2.18	3.01	2.07	4.00	2.29	2.71
868308	65.99	24.35	3.48	1.83	1.37	4.88	2.00	2.71
429906	5.35	1.98	1.98	3.10	2.51	2.18	3.77	2.71
83011	32.97	12.18	2.39	2.09	2.06	3.35	3.64	2.71
838611	52.82	19.51	4.00	0.88	1.23	4.84	2.60	2.71
783697	3.19	1.18	2.47	3.13	3.04	2.62	2.26	2.70
268178	28.02	10.37	1.92	3.23	2.46	3.05	2.85	2.70
460487	24.08	8.92	0.68	0.85	5.68	0.08	6.21	2.70
418053	7.57	2.80	3.25	1.89	2.22	3.44	2.70	2.70
884894	41.65	15.44	3.11	2.60	1.96	3.14	2.68	2.70
824066	4.08	1.51	2.48	2.33	1.91	5.02	1.74	2.70
950450	1.77	0.66	1.96	2.45	3.15	3.54	2.39	2.70
50344	0.85	0.32	1.00	1.06	0.91	9.45	1.06	2.70
42076		2.12	2.14	3.48	2.38	2.44	3.04	2.69
68636		1.80	2.78	3.27	2.43	3.37	1.62	2.69
26185	1 1	0.43	1.89	2.57	3.38	3.81	1.78	2.69
278687		4.88	2.06	4.03	2.38	1.89	3.07	2.69
391984	4.56	1.70	3.22	3.44	1.67	2.03	3.06	2.68
279785	: :	0.34	1.97	1.94	1.70	5.89	1.92	2.68
897199		5.51	3.48	2.55	2.61	3.07	1.71	2.68
263697		1.58	3.06	3.23	4.13	1.12	1.87	2.68
770983	E i	2.70	1.30	5.12	2.60	2.36	2.03	2.68
505344	•		3.69	2.24	1.96	3.13	2.38	2.68
253865			5.25	3.14	1.49	1.55	1.93	2.67
752690	4	1 1	3.98	2.14	0.98	4.23	2.02	2.67
857681			3.42	2.88	1.57	3.42	2.07	2.67
1049033	•		2.50	3.09	3.57	3.21	0.99	2.67
743041			3.55	3.05	3.00	2.00	1.76	2.67
461761			2.63	1.55	4.32	3.11	1.72	2.67
51921	L .		5.04	1.90	2.84	1.30	2.26	2.67
1471829 307882	1		2.97 3.13	2.21 1.77	1.62	5.24	1.30 2.12	2.67 2.67
74308	L		3.13 4.46	3.27	2.17	4.14 2.05	2.12	2.67
74308 251452			4.46 1.29		1.20	2.05	1.24	
79573			3.12	2.18	2.99	. 5.63		2.67 2.66
83847	l .		3.12 4.43	3.28	1.60 2.84	3.79 1.08	1.52 3.16	2.66
36132	1		4.43 1.66	1.79 1.41	2.84	6.37	1.76	2.66
52986			2.44	1.41 2.09	2.10 3.05	3.43	2.29	2.66
32760	1 3./.	7 2.10	2.44	2.09	3.03	3.43	2.29	2.00

									0.66
	486175	10.39	3.91	1.74	7.19	2.53	0.92	0.91	2.66
	307029	56.38	21.22	3.05	1.98	1.59	4.58	2.09	2.66
	243321	2.79	1.05	2.81	1.53	1.43	1.74	5.74	2.65
	590398	12.27	4.63	4.11	2.22	2.03	3.29	1.58	2.65
	1475028	249.73	94.34	2.17	1.81	1.81	4.31	3.14	2.65
	595604	9.41	3.56	1.72	1.96	4.39	3.06	2.10	2.65
	188151	3.25	1.23	1.76	5.51	1.64	2.21	2.09	2.64
	859832	2.57	0.97	1.88	3.53	2.79	3.19	1.81	2.64
	42313	1.33	0.51	3.37	2.77	2.16	3.35	1.55	2.64
	282144	30.16	11.43	1.47	2.73	2.45	5.25	1.30	2.64
	32887	3.04	1.16	1.78	3.54	3.59	2.10	2.17	2.64
	137254	1.35	0.51	3.82	2.00	4.43	1.38	1.55	2.64
	234036	4.77	1.81	1.87	3.76	2.13	3.14	2.28	2.64
	206785	6.05	2.30	4.91	3.70	1.64	1.42	1.50	2.63
	244202	9.25	3.51	1.51	2.29	3.36	1.92	4.09	2.63
	827144	2.52	0.96	1.88	3.18	2.21	1.64	4.24	2.63
	838504	41.97	15.95	1.67	3.37	2.82	3.53	1.77	2.63
	376764	2.58	0.98	1.35	2.16	3.58	3.03	3.03	2.63
	744933	2.15	0.82	0.98	1.12	0.87	8.87	1.30	2.63
	161456	13.35	5.08	0.92	0.15	10.09	0.13	1.83	2.63
4	564846	1.90	0.72	3.47	1.34	2.53	2.17	3.62	2.63
	815072	3.49	1.33	5.03	2.36	1.34	2.07	2.32	2.62
	1599489	5.51	2.10	3.32	3.51	1.94	2.95	1.40	2.62
Controlled Controlled	80292	2.18	0.83	2.30	3.64	1.47	3.07	2.62	2.62
E 2 2	262865	30.48	11.64	3.50	2.06	4.94	0.99	1.61	2.62
200 M	590390	6.42	2.45	4.07	2.94	1.83	3.34	0.92	2.62
	610011	34.63	13.23	3.40	2.31	2.21	1.95	3.23	2.62
1	855487	8.56	3.27	2.21	3.52	3.16	1.26	2.94	2.62
271 271	503097	7.27	2.78	1.87	1.80	7.01	1.54	0.86	2.62
100000 1000000	825719		0.29	3.01	3.07	3.22	2.78	0.99	2.62
	282720		3.18	3.44	1.29	2.89	2.24	3.20	2.61
STATE OF THE PARTY	767798		2.18	2.43	1.90	2.16	3.46	3.13	2.61
	784129		2.50	1.21	2.18	5.46	2.80	1.41	2.61
77777	23433			2.32	2.24	3.25	2.18	3.04	2.61 2.61
	73897			1.21	2.29	7.41	1.31	0.82	2.60
,	4141	1 112.58		3.28	2.76	1.54	3.09	2.35	
	78426		1 L	1.81	3.29	2.15	3.61	2.13	2.60 2.60
	81324			3.66	1.96	4.73	1.34	1.29	2.60
	84053	1	1 1	3.14	3.14	1.61	3.33	1.76	2.59
-	81175	1	3 I	3.15	2.92	3.96	1.62	1.32 1.74	2.59
	50236	3		3.58	2.45	2.17	3.01	3.44	2.59
	74494	I .		3.76	1.13	2.12	2.49	5.42	2.59
	85563			1.55	1.65	3.18	1.13	1.65	2.58
	7261	4	1 1	3.07	3.09	2.14	2.94	1.65	2.58
	43610		1 1	2.43	2.14	3.01	3.83 3.12	3.67	2.58
	4873			2.52	2.05	1.52	1.52	5.87	2.57
	8141	· ·		1.83	1.77	1.85	5.65	1.19	2.57
	4353		1 1	1.68	2.24	2.09	3.63 4.15	2.62	2.57
	8385		1 1	1.53	3.21	1.33 1.89	2.17	3.08	2.57
	1376				2.54		3.58	2.07	2.57
	412	07 2.0	5 0.80	1.72	2.29	3.16	٥د.د	2.07	2.31

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	1558642	3.11	1.21	3.27	2.26	1.72	4.08	1.50	2.56
	432194	10.34	4.04	1.58	3.07	2.91	1.91	3.34	2.56
	232628	4.02	1.58	4.00	1.88	4.50	1.05	1.32	2.55
	951022	3.02	1.18	3.66	2.02	3.80	1.61	1.67	2.55
	809824	2.66	1.05	1.68	2.11	2.41	3.25	3.24	2.54
	897906	10.34	4.08	1.96	2.02	4.28	0.64	3.78	2.54
	809722	50.59	19.96	0.71	0.97	3.43	3.59	3.97	2.54
	138865	5.42	2.14	1.16	3.20	2.53	3.12	2.65	2.53
	825012	4.46	1.76	3.39	2.40	2.05	3.22	1.61	2.53
	1472689	5.15	2.03	1.97	0.59	1.72	7.13	1.27	2.53
	322537	3.18	1.26	1.26	6.07	2.04	0.89	2.40	2.53
	291974	57.24	22.63	3.29	1.87	1.09	4.48	1.91	2.53
	307231	2.04	0.81	1.95	1.61	1.41	6.48	1.19	2.53
	1638479	3.27	1.30	2.08	3.88	3.23	1.78	1.67	2.53
	415415	9.68	3.83	3.25	2.43	1.49	3.21	2.26	2.53
	251417	0.69	0.27	3.59	3.13	1.67	1.79	2.46	2.52
	452936	5.15	2.04	3.47	2.84	3.54	1.65	1.11	2.52
	488301	0.97	0.38	2.10	1.37	1.81	5.57	1.76	2.52
	430235	3.43	1.36	4.46	3.50	1.54	1.77	1.34	2.52
	784085	3.93	1.56	3.14	2.04	3.39	2.55	1.49	2.52
	741988	1.74	0.69	1.90	3.05	2.20	3.46	1.99	2.52
:	627688	7.37	2.93	1.52	1.61	6.90	1.77	0.79	2.52
	825859	1.64	0.65	1.30	2.26	2.23	3.48	3.31	2.52
	884892	8.02	3.19	1.99	2.90	3.01	3.04	1.63	2.52
	277660	14.16	5.65	1.86	2.15	2.18	3.13	3.22	2.51
	76605	2.48	0.99	4.39	3.21	1.27	2.11	1.53	2.50
	220069	4.81	1.92	1.69	3.29	3.04	1.86	2.61	2.50
	460798	1.30	0.52	1.95	1.79	3.77	1.74	3.24	2.50
	247546	11.16	4.48	1.50	1.33	1.37	2.54	5.72	2.49
	814080	8.19	3.29	1.76	1.45	2.40	1.52	5.32	2.49
	1422794	2.62	1.05	0.95	1.99	3.82	3.31	2.38	2.49
	51981	19.62	7.89	3.06	1.82	1.65	2.42	3.50	2.49
	325606	7.30	2.93	3.10	2.79	1.65	3.28	1.62	2.49
	277596	2.91	1.17	1.17	2.54	1.50	6.20	1.01	2.49
	788205	13.21	5.32	3.46	2.10	1.43	1.78	3.65	2.48
	796527	1.35	0.55	2.22	1.81	1.52	5.45	1.39	2.48
	827132	6.63	2.68	2.47	1.83	1.62	3.44	3.03	2.48
	842863	3 11.11	4.49	3.25	2.37	1.47	2.24	3.06	2.48
	29964	1.12	0.45	1.27	1.88	1.58	4.63	3.01	2.47
	13798	4 1.63	0.66	1.72	2.25	1.34	6.05	1.01	2.47
	76734		1.82	1.80	1.32	6.76	1.74	0.73	2.47
	149296	7 3.68		2.19	1.49	6.63	1.16	0.89	2.47
	34272	1		3.30	1.93	5.09	0.69	1.30	2.46
	7600			4.18	3.61	1.37	1.59	1.56	2.46
	82385	4		2.69	1.51	1.29	5.06	1.76	2.46
	74490	1		3.34	2.72	1.92	3.17	1.16	2.46
	39953	1	1 1	1.68	2.54	1.91	3.15	3.01	2.46
	95046	1	5 E	1.50	1.01	7.67	1.11	0.97	2.45
	48634	1		1.85	3.58	1.43	3.51	1.87	2.45
	86686		1 1	4.47	3.47	1.46	1.44	1.40	2.45
	88454	156.92	64.25	2.76	1.70	1.46	3.06	3.23	2.44

897448	31.17	12.76	3.09	2.67	1.77	3.01	1.67	2.44
950989	2.48	1.02	4.36	3.15	1.27	1.86	1.55	2.44
1031583	38.01	15.59	0.37	1.52	1.75	6.11	2.45	2.44
381036	3.54	1.46	2.61	1.83	5.63	1.30	0.81	2.43
109316	12.35	5.08	1.34	0.40	2.84	0.34	7.26	2.43
448117	3.91	1.61	3.94	1.97	3.08	1.50	1.68	2.43
156045	2.01	0.83	3.16	3.14	1.81	2.33	1.71	2.43
884333	4.10	1.69	2.33	3.28	3.72	1.30	1.51	2.43
306771	3.61	1.49	3.02	3.83	1.70	1.84	1.73	2.42
430465	5.55	2.30	0.96	0.93	1.81	7.45	0.92	2.42
123755	5.06	2.10	3.75	1.13	1.03	1.10	5.05	2.41
42803	2.96	1.23	0.60	2.20	1.73	5.20	2.32	2.41
460164	0.80	0.33	2.54	3.13	2.35	3.08	0.95	2.41
1574594	3.08	1.28	4.44	3.10	1.44	1.96	1.07	2.40
141361	1.28	0.54	1.80	0.83	1.31	6.11	1.94	2.40
855336	5.69	2.37	1.65	3.16	2.36	3.39	1.43	2.40
868396	6.77	2.83	5.03	3.73	0.87	1.05	1.28	2.39
627053	11.04	4.64	1.29	2.07	5.16	2.01	1.37	2.38
815772	1.44	0.61	3.35	1.86	3.10	1.62	1.96	2.38
897547	15.69	6.62	3.93	3.84	2.46	0.78	0.84	2.37
202154	29.46	12.46	0.97	2.61	0.99	5.43	1.83	2.37
511107	7.14	3.02	0.73	1.70	3.61	3.54	2.25	2.36
810408	9.85	4.18	1.91	1.77	1.69	3.08	3.34	2.36
757222	5.14	2.18	2.47	6.61	1.12	0.68	0.91	2.36
810734	2.46	1.05	3.06	1.80	1.50	4.04	1.32	2.35
1493390	3.57	1.53	3.30	1.07	2.47	3.35	1.54	2.34
284865	2.87	1.23	3.64	3.21	1.66	1.49	1.70	2.34
488033	3.14	1.34	2.41	3.80	1.46	3.21	0.82	2.34
37665	1.01	0.43	0.55	3.88	2.77	3.16	1.33	2.33
138861	1.72	0.74	1.34	3.12	1.63	3.22	2.35	2.33
272049	6.87	2.95	3.26	1.35	1.69	4.65	0.71	2.33
427812	9.94	4.27	2.58	1.48	5.45	1.15	0.97	2.33
809995	2.80	1.21	3.69	1.55	2.04	3.10	1.24	2.32
449275	3.81	1.64	5.51	1.68	1.24	2.01	1.16	2.32
731337	0.75	0.32	3.30	3.09	1.35	1.91	1.95	2.32
854570	4.11	1.78	4.79	3.45	0.94	1.54	0.83	2.31
767988	3.64	1.58	1.57	3.20	3.21	1.89	1.63	2.30
455271	1.64	0.72	5.02	2.60	1.17	1.48	1.22	2.30
130136	0.29	0.13	1.17	1.08	0.84	0.80	7.57	2.29
757191	6.08	2.66	1.09	0.88	5.46	2.03	1.99	2.29
266500	7.14	3.12	1.54	3.23	1.95	3.65	1.08	2.29
725503	4.55	1.99	3.35	2.31	1.33	3.39	1.03	2.28
1555523	1.21		1.80	6.52	1.30	0.93	0.86	2.28
629498	3.73	1.64	4.45	3.42	1.09	1.19	1.23	2.28
781401		1 1	1.40	1.34	2.90	0.55	5.16	2.27
632137	0.80		4.14	1.33	3.68	1.07	1.12	2.27
293830	· ·	1 1	1.87	1.37	5.66	1.25	1.18	2.26
50562	1		6.49	1.87	0.28	0.43	2.21	2.26
243784	,	1 1	3.68	3.50	1.79	1.35	0.94	2.25
20916	1		5.12	1.17	2.88	0.92	1.17	2.25
25172	7 4.88	2.17	1.20	3.39	1.73	3.26	1.65	2.25

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284545	1.10	0.49	1.06	3.27	0.97	4.66	1.26	2.24
504290	50.88	22.72	1.27	1.74	5.93	1.07	1.18	2.24
71672	4.77	2.14	3.51	1.66	3.03	1.97	0.99	2.23
279394	3.18	1.42	1.37	1.39	5.66	0.95	1.78	2.23
309864	2.49	1.12	8.06	0.69	0.90	0.79	0.67	2.22
291394	6.46	2.92	1.70	1.25	5.58	1.18	1.37	2.22
289645	0.92	0.42	1.62	1.02	1.30	5.62	1.50	2.21
460218	2.10	0.95	1.71	1.76	3.37	3.35	0.86	2.21
856174	0.25	0.11	3.70	4.65	0.88	0.91	0.92	2.21
767784	8.86	4.05	5.50	1.03	1.73	1.72	0.97	2.19
826459	12.36	5.65	0.94	3.16	1.94	3.18	1.72	2.19
141336	2.12	0.97	0.78	0.91	7.64	0.64	0.94	2.18
122239	6.50	2.98	4.86	3.17	0.88	0.92	1.06	2.18
840698	7.45	3.42	3.42	3.27	1.39	1.58	1.22	2.18
303139	2.17	1.00	0.98	0.85	2.90	5.31	0.82	2.17
277761	2.43	1.12	3.26	1.81	1.46	3.22	1.08	2.17
471641	4.79	2.22	0.68	4.45	0.89	4.15	0.64	2.16
470114	3.09	1.44	2.04	3.03	1.58	3.16	0.97	2.15
33045	0.63	0.29	5.38	1.08	1.25	1.95	1.09	2.15
84078	0.34	0.16	0.79	0.75	3.03	4.97	1.17	2.14
111389	2.59	1.21	3.22	1.26	1.20	3.23	1.78	2.14
1536991	10.79	5.05	3.61	1.25	3.84	1.06	0.92	2.14
770935	86.21	40.44	0.25	2.50	1.55	3.29	3.08	2.13
281191	7.36	3.45	3.30	1.86	3.23	0.81	1.46	2.13
149895	: ;	2.84	1.33	1.44	6.25	0.94	0.69	2.13
648046	1 1	7.32	0.56	0.94	3.27	2.53	3.34	2.13
855236	1 1	1.30	1.11	1.65	1.52	5.25	1.06	2.12
740780		2.25	1.78	6.08	0.90	0.38	1.35	2.10
844768		3.08	4.79	3.24	0.75	0.87	0.85	2.10
811097	1 1	1.23	5.78	1.43	0.76	1.89	0.62	2.10
731255		0.98	1.12	1.83	1.35	5.14	1.01	2.09
950607	1	9.50	2.04	1.14	5.08	1.05	1.11	2.08 2.08
1472150		1	3.00	1.41	1.13	3.73	1.12 1.07	2.08
52338			3.16	1.49	4.05	0.57 3.43	0.88	2.05
781061			0.76	1.88	3.32 0.88	2.54	0.68	2.05
1613940			3.02	3.14		0.92	0.85	2.03
160471			1.03	1.41	5.95 0.96	5.73	0.57	2.02
15147			0.96	1.86	0.90	3.64	0.86	2.01
56344	L.		1.64	3.21 0.82	5.31	1.07	0.95	2.01
82716			1.91 4.42	3.23	0.81	0.67	0.89	2.00
84578 74357	E.		4.42	3.23	0.89	0.69	0.88	2.00
43159	1		1.67	0.89	5.75	0.72	0.97	2.00
74094	B .	1 1	6.11	1.20	0.62	0.67	1.29	1.98
34368			1.39	1.27	5.47	0.79	0.88	1.96
7505	1		3.00	1.01	1.04	3.21	1.50	1.95
78251	E .	1 1	1.16	0.84	5.53	1.41	0.75	1.94
36681			5.26	1.35	1.25	0.46	1.38	1.94
13291	l l	1 1	3.07	1.21	3.40	0.74	1.22	1.93
28289		1 :	1.03	1.04	5.87	0.90	0.81	1.93
5121		1 1	1.08	0.94	5.62	1.08	0.80	1.90
J.21	1						•	

291706	2.41	1.27	1.34	1.06	0.98	0.91	5.20	1.90
50114	0.83	0.44	0.97	1.06	0.39	3.57	3.45	1.89
450464	3.97	2.13	3.23	1.41	3.18	0.56	0.95	1.86
626462	5.86	3.24	3.91	3.03	0.70	0.59	0.81	1.81
867717	5.61	3.15	3.27	3.01	0.87	0.68	1.08	1.78
462953	3.21	1.90	0.67	5.53	1.05	0.63	0.57	1.69
41287	0.72	0.45	0.73	0.95	0.53	5.14	0.61	1.59
122428	2.03	1.29	5.48	0.55	0.63	0.63	0.58	1.57
1227201	2.03		• • • •				•	

Table 1-2

	Clone	Aver of	Aver. of	MPM-115/	MPM-116/	MPM-117/	MPM-118/	MPM-119 /	Aver.
			Normals	Aver	Aver	Aver		Aver Normal	Fold
•	788180	25.18	0.81	62.98	3.35	9.48	55.67	24.72	31.24
	646037	24.42	1.02	38.80	10.71	20.96	13.10	35.99	23.91
	1034473	33.61	1.51	39.30	2.42	16.40	18.87	34.35	22.27
	281003	128.77	6.30	25.82	30.44	5.57	24.33	15.96	20.43
	773479	12.32	0.65	24.85	24.44	9.41	16.91	18.67	18.85
	823940	22.39	1.32	15.54	10.31	45.60	5.40	8.09	16.99
	281039	14.05	0.94	1.73	6.90	1.46	63.60	1.27	14.99
	46166	13.13	0.93	2.81	31.75	16.71	15.27	4.05	14.12
	42415	6.86	0.51	4.67	6.71	15.78	1.60	38.87	13.52
	291255	15.20	1.35	10.61	10.39	17.35	9.54	8.56	11.29
	308682	8.99	0.87	8.35	12.44	6.96	9.39	14.72	10.37
	450049	18.37	1.83	8.45	6.05	7.47	5.21	23.07	10.05
	283312	63.94	6.63	11.46	5.48	5.23	11.97	14.09	9.65
	432194	10.34	1.11	5.77	11.20	10.61	6.96	12.20	9.35
	35626	9.42	1.09	6.10	9.04	16.67	3.16	8.40	8.67
	222157	8.80	1.02	13.68	8.34	7.21	3.07	11.05	8.67
	814053	14.66	1.70	8.20	1.30	6.73	19.59	7.22	8.61
	811867	24.57	2.97	9.33	7.93	6.31	11.68	6.18	8.28
	35058		1.89	3.32	10.65	5.05	16.86	5.29	8.24
	971279		6.63	8.84	8.00	7.58	8.74	7.81	8.20
- TOT-	293635	1	3.44	5.13	10.21	7.80	7.55	7.78	7.69
	32160	1	3.59	6.47	7.13	8.59	6.37	8.77	7.47
uj	754280	1	3.20	15.06	7.80	8.20	3.02	3.20	7.46
	282089			1.86	11.16	1.40	21.82	0.82	7.41
	745347	1	•	3.74	8.94		8.80	6.91	7.39
TOTAL STREET	67401		•		1.47	30.35	2.21	1.11	7.28
	293651	13.35	1.84	3.60	7.75	11.33	6.53	7.12	7.26
	784168	53.04	7.37	2.50	6.64	8.80	8.22	9.84	7.20
	811603	19.62	2.74	8.80	6.79	5.12	7.74		1
	486035	24.75	3.47	5.79	11.37	4.91			1
	809603	49.03	6.94	5.63	5.54	6.31	7.30		1
	1048810	20.15	3.04	7.11	10.78	3 4.14			
IJ F	756600	47.06	5 7.13	6.98	3.61	1 3.71			I.
	77230	4 113.14	18.10	7.98	4.62	2 4.60			1
	163171	3 5.68	0.93						1
	83836	4							i i
	75743	5 365.7							
	155652			1					
	45030		I .						1
	79616	B .		L .					
	103473	1							
	4930		L L						
·	29251								1
	81022								
	20702			1					•
	41570								
	44911	· ·	1						
	53102	1		E .					1
	36275								1
	50988								4
	19190	•		(1
	8778:	35 23.0	06 4.3	14 7.4	18 3.0	06 4.3	32 5.1	71 6.0	5.32

					· · •			
771317	9.44	1.78	4.49	3.90	5.78	4.69		5.31
549101	12.81	2.43	7.81	3.40	3.06	5.03		5.26
84955	5.26	1.01	8.05	6.43	6.35	2.52		5.21
461670	8.90	1.72	3.21	6.57	7.68	3.75		5.19
197525	9.52	1.85	2.67	8.08	4.07	6.57	3	5.16
730739	36.49	7.10	4.49	4.01	10.50	3.62		5.14
240694	4.53	0.90	4.65	7.82	3.10	3.41		5.01
593185	28.06	5.61	1.33	3.36	3.94	10.92		5.00
212165	41.96	8.42	7.66	4.81	6.54	2.71		4.99
268211	3.43	0.70	4.14	3.18	3.13	2.23		4.93
233419	5.55	1.14	3.13	6.59	6.63	2.94		4.87
296788	3.27	0.67	4.84	4.90	7.08	2.63	4.79	4.85
810558	14.49	3.00	3.18	4.47	4.01	4.93	7.57	4.83
264895	7.86	1.63	3.07	6.53	4.76	3.25	6.52	4.82
51773	11.48	2.40	5.99	3.55	3.22	7.62	3.50	4.77
191603	22.14	4.64	6.96	6.89	2.33	3.85	3.81	4.77
417908	12.10	2.57	9.22	5.41	4.37	1.99	2.57 4.92	4.71 4.69
50188	10.75	2.29	5.42	4.91	4.42	3.76	2.86	4.68
284701	33.05	7.05	3.72	4.93	2.53	9.40 4.54	1.77	4.67
416436	11.08	2.37	7.73	5.37	3.95	2.65	5.05	4.61
321470	7.74	1.68	4.78	6.01	4.56 2.44	2.63 3.97	2.28	4.58
449275 263697	3.81 4.23	0.83 0.93	10.88 5.22	3.32 5.51	7.03	1.91	3.18	4.57
266106	20.68	4.53	3.22 4.04	3.31 4.70	9.22	2.05	2.82	4.57
194364	5.44	1.19	2.81	3.35	9.83	4.15	2.64	4.56
259374	10.51	2.33	4.05	2.96	1.50	8.52	5.48	4.50
877641	67.93	15.10	8.40	2.61	1.99	5.49	3.99	4.50
283173	1	3.97	10.94	2.48	2.73	4.55	1.75	4.49
868282	i	2.01	4.17	4.41	5.09	4.45	4.28	4.48
770066		2.60	1.11	1.71	4.03	13.55	1.61	4.40
950688		11.99	5.18	2.58	3.06	4.94	6.20	4.39
490178	1	5.63	2.55	2.07	13.03	1.58	2.57	4.36
30428	•	0.99	0.90	2.03	2.95	11.84	4.01	4.35
210405	E	7.84	3.39	3.97	5.43	4.21	4.69	4.34
795907	1	1.25	1.95	2.24	4.60	0.83	11.91	4.31
47542	I	1.88	6.90	3.68	2.63	4.23	4.04	4.30
297439		•	4.18	7.63	3.83	1.95	3.86	4.29
1472689	5.15	1.21	3.31	0.99	2.88	11.98	2.13	4.26
89776	7 2.07	0.49	3.67	3.81	4.12	3.26	6.41	4.25
28989	1 7.82	1.84	4.50	2.99	9.38	2.41	1.95	4.25
85396	8 12.10	2.86	5.03	5.67	7.48	1.76	1.23	4.23
84078	8 23.03	5.46	4.18	3.11	1.97	7.77	4.07	4.22
85644				2.15	2.21	4.88	2.79	4.22
34688	1		•	5.10	1.70	6.26	5.45	4.21
32170	1		B .	0.98	9.88	1.19	3.34	4.20
85574	1			3.70	3.56	3.56	4.54	1
103158		1		2.62	3.01	10.51	4.21	
78591			1	3.95	7.94	2.07	2.99	1
74615		1	1	4.05	7.60	1.47	0.48	
84536		1	•	3.53	2.38	5.97	4.42	•
37724	1		L	2.77	11.38	1.14	1.90	ł
12902			1	0.75	2.32	2.43	7.90	t .
15020				5.14	4.58	3.46	3.56 3.94	
82526	07 11.9	6 2.93	2 3.12	3.72	4.18	5.49	3.94	4.09

1030855	14.86	3.64	1.14	4.82	2.58	7.86	****	4.08
878127	3.53	0.87	2.76	4.48	6.05	4.44		4.07
138444	4.04	1.00	3.68	7.21	2.94	4.04		4.06
768395	14.29	3.54	5.36	4.80	4.35	3.14	1	4.03
788541	12.11	3.01	2.87	3.90	7.56	1.94		4.03
1468260	4.53	1.13	7.56	0.78	4.31	2.33		4.03
787938	5.09	1.27	1.46	4.56	6.98	5.35	1.72	4.01
258300	18.15	4.53	2.52	4.17	7.79	0.73	4.84	4.01
208082	7.90	1.98	3.29	6.85	3.52	3.30	3.00	3.99
785910	4.01	1.01	3.80	2.26	9.32	1.33	3.21	3.98
287728	2.76	0.70	0.75	5.13	1.25	11.82	0.89	3.97
687297	12.18	3.12	4.57	4.32	4.27	2.73	3.63	3.90
741067	64.30	16.50	0.91	4.38	2.37	7.31	4.51	3.90
1031598	5.18	1.34	0.69	2.70	2.63	9.79	3.58	3.88
366815	71.61	18.49	10.51	2.69	2.49	0.91	2.76	3.87
323474	1.25	0.32	6.77	4.12	2.60	2.99	2.88	3.87
288695	3.41	0.88	4.88	4.64	5.25	3.06	1.48	3.86
204111	3.38	0.87	3.40	4.11	3.20	3.52	5.07	3.86
242952	16.94	4.39	4.50	3.65	3.98	4.16	2.99	3.86
1631747	5.09	1.33	5.74	6.76	2.76	2.44	1.49	3.84
48286	2.34	0.61	3.34	2.74	4.95	3.47	4.67	3.83
745503	42.62	11.14	2.97	4.02	3.39	3.08	5.68	3.83
744846	2.69	0.70	4.56	5.57	3.31	4.45	1.24	3.83
454771	2.69	0.70	7.92	2.22	2.30	1.94	4.70	3.82
233464	14.24	3.73	2.79	2.89	5.83	3.29	4.26	3.81
250869	123.92	32.54	10.99	2.78	2.27	1.03	1.97	3.81
842925	4.61	1.21	3.52	3.54	4.61	4.62	2.67	3.79
843094	49.18	13.00	4.51	3.53	3.71	3.00	4.17	3.78
587415	12.98	3.44	6.76	3.20	6.42	1.08	1.39	3.77
33715		0.37	2.84	3.84	4.39	3.91	3.66	3.73
75044	ł	0.93	4.59	3.85	2.61	3.71	3.84	3.72
1467207	1	0.99	2.51	4.57	4.41	2.93	4.17	3.72
366132	1	1.68	4.94	2.88	2.87	4.12	3.77	3.72
813973	1	0.88	3.24	3.78	4.03	3.23	4.23	3.70
814017	1	23.42	5.13	5.73	3.05	1.59	2.90	3.68
785616		3.12	4.63	4.47	3.42	3.47	2.37	3.67
42803	1	0.81	0.91	3.34	2.63	7.89	3.52	· 3.66
488964	1	2.13	11.81	2.73	1.04	1.76	0.88	3.64
151055	3		3.65	6.52	1.80	2.91	3.31	3.64
784258		1	4.79	2.92	5.87	1.93	2.67	3.64
81331	1	1	1.10	2.95	0.88	12.05	1.18	3.63
897042			6.51	3.85	2.87	3.80	1.14	3.63
771048	3		4.89	3.20	3.32	2.85	3.91	3.63
795265			1.26	2.63	2.84	9.65	1.76	3.63
753253			2.88	3.17	3.38	4.62	4.10 1.38	3.63 3.62
43046		•	1.44	1.39	2.71	11.18	2.08	1
42967	1		L .	2.41	5.14	4.64	2.08	3.61 3.61
4738			1	3.72	4.97	3.42 6.29	2.98	3.60
34227			t .	1.46	4.16 3.05	11.90	0.65	3.59
71295		1	8	1.53	3.05 3.91	4.02	1.91	3.59
24554	•	•		6.79	3.91	4.02 6.04	2.26	
147598 35885	1		t .	2.40 3.37	2.69	1.80	3.82	1
4532		•	1	3.39	3.66	4.62	3.78	
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	Table 1-2											
950410	3.18	0.89	1.81	7.55	3.00	2.68	2.76	3.56				
1592006	5.30	1.51	4.10	3.01	3.34	3.68		3.52				
741474	4.93	1.40	2.90	2.23	3.62	5.57	3.24	3.51				
292042	9.39	2.67	3.33	4.22	4.13	3.14	2.73	3.51				
280217	20.52	5.86	4.76	0.46	1.34	10.23	0.72	3.50				
42636	1.11	0.32	2.78	3.16	4.86	2.57	4.13	3.50				
550141	78.20	22.40	5.62	2.32	2.57	3.16	3.79	3.49				
159608	14.43	4.14	6.82	1.07	1.06	5.09	3.39	3.49				
383185	12.04	3.46	6.25	2.54	6.69	0.98	0.94	3.48				
491184	6.35	1.83	4.07	1.43	10.01	0.85	1.00	3.47				
786525	1.46	0.42	2.15	1.29	2.53	10.04	1.31	3.46				
305538	11.44	3.31	4.78	4.20	1.98	1.85	4.49	3.46				
415437	3.52	1.02	2.69	3.45	3.10	4.16	3.88	3.46				
259350	22.84	6.64	2.91	2.83	2.97	3.77	4.71	3.44				
950709	4.17	1.22	5.93	2.37	2.11	4.13	2.61	3.43				
141505	6.38	1.86	2.88	4.20	3.06	2.08	4.90	3.42				
109200	3.17	0.93	5.89	2.59	3.19	2.96	2.46	3.42				
1390584	6.54	1.91	1.98	3.12	3.26	8.11	0.63	3.42				
823819	11.96	3.51	4.50	4.09	2.11	2.94	3.43	3.41				
272750	20.57	6.04	3.75	5.31	2.77	2.65	2.54	3.40				
246869	4.68	1.38	4.93	2.40	3.61	2.07	3.98	3.40				
784253	11.76	3.47	2.81	4.36	3.28	2.90	3.61	3.39				
490766	3.48	1.03	4.47	3.60	2.64	2.91	3.27	3.38				
356863	3.18	0.94	4.17	3.83	2.02	3.24	3.63	3.38				
837904	103.13	30.63	2.21	2.23	2.11	4.52	5.77	3.37				
462468	3.24	0.96	3.67	4.25	3.86	1.95	3.06	3.36				
121997	5.10	1.52	4.66	2.43	5.54	1.60	2.55	3.36				
700900	9.94	3.00	3.26	2.88	3.29	3.75	3.40 3.40	3.31 3.31				
26616 1589973	1.94	0.59	4.02 2.96	2.79 3.72	3.28 2.30	3.05 5.22	2.31	3.30				
392678	5.17 34.95	1.57 10.61	2.96	3.72 2.72	3.38	3.80	4.32	3.29				
586725	2.32	0.70	3.69	2.72	2.18	3.31	4.64	3.29				
121489		10.55	3.98	4.08	3.24	2.67	2.47	3.29				
503851	E .	5.54	1.87	3.27	5.43	2.90	2.90	3.28				
854874	ł	0.68	2.50	3.02	5.99	2.47	2.32	3.26				
877832	1	13.65	3.45	3.31	2.92	3.09	3.50	3.25				
253009	•	1	1.83	3.37	1.51	5.38	4.13	3.24				
824723	l .	•	2.37	2.55	3.40	5.30	2.52	3.23				
755409			3.30	3.12	4.05	3.18	2.47	3.22				
506564		l .	4.44	2.86	3.21	1.98	3.55	3.21				
109153	•		4.21	2.85	3.08	2.33	3.55	3.20				
213682			1	2.78	2.14	7.22	3.16	3.19				
133717	9.46	2.97	2.19	2.22	3.36	2.66	5.49	3.18				
361659	13.43	4.23	1.54	3.83	4.60	3.93	1.99	3.18				
23262	8 4.02	1.27	4.95	2.33	5.56	1.30	1.63					
28303	9.61	3.05	3.07	5.05	2.86	2.31	2.46	1				
49023			L L	3.48	3.06	1.82	3.75	4				
78136			1	2.65	1.25	2.22	6.10					
6631		4		2.02	1.77	4.59	0.44					
35978			a to the second	5.41	2.09	3.84	2.45					
56480				1.38	8.87	1.73	0.67	1				
58759		•		2.46	3.02	6.40	2.98	L				
3988	1		•	2.92	4.86	4.97	1.87					
4557	8 5.0	5 1.63	4.85	2.27	2.38	3.60	2.39	3.10				

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743749	5.10	1.65	4.98	3.05	2.98	3.05		3.10
839081	16.60	5.37	5.25	5.49	0.97	2.62		3.09
109316	12.35	4.01	1.69	0.50	3.59	0.43		3.08
358083	2.45	0.80	3.44	2.58	4.41	1.95		3.07
130773	3.71	1.21	2.86	3.98	3.19	2.42		3.07
610011	34.63	11.27	3.99	2.71	2.59	2.28	1	3.07
429574	3.77	1.23	2.74	4.01	3.30	1.01	li li	3.06
167205	7.85	2.56	2.90	5.74	3.48	1.88		3.06
825718	4.03	1.32	2.59	2.65	4.00	3.21	2.84	3.06
42659	10.21	3.36	3.63	3.18	3.06	2.09	3.23	3.04
1631682	1.61	0.53	4.08	2.09	3.06	2.23	3.71	3.03
843234	7.02	2.32	3.12	3.14	2.40	2.69	3.80	3.03
290642	7.76	2.56	4.18	2.71	2.32	3.00	2.91	3.02
51700	3.66	1.21	3.02	2.75	3.37	2.08	3.87	3.02
214006	4.67	1.55	6.05	4.62	1.41	2.11	0.89	3.02
431286	13.81	4.59	2.40	3.23	3.52	2.10	3.80	3.01
781454	9.75	3.24	2.57	1.19	9.17	1.13	0.98	3.01
433253	7.29	2.43	2.31	2.35	2.05	5.26	3.03	3.00
359661	26.24	8.79	2.02	4.95	1.82	3.95	2.19	2.99
811953	17.84	5.98	3.25	3.61	3.69	1.97	2.41	2.98
196345	122.93	41.22	1.48	2.47	2.71	3.86	4.39	2.98
950473	9.36	3.14	2.36	3.35	2.97	2.65	3.56	2.98
46294	1.03	0.35	1.45	2.41	1.44	5.75	3.83	2.97
37980	0.84	0.29	1.32	2.54	1.61	7.51	1.81	2.96
512116	4.59	1.55	1.67	1.46	8.72	1.68	1.24	2.96
137638	1	1.71	3.63	2.92	2.17	2.49	3.54	2.95
810753	•	1.68	3.05	2.91	3.48	2.96	2.35	2.95 2.95
592403	1	0.16	0.70	2.31	2.27	6.90 2.44	2.56 2.71	2.95 2.95
273546	1	3.09	4.02	3.24	2.32	2.44 0.62	6.02	2.94
238821	1	2.22	3.78	1.07 4.26	3.22 4.44	1.20	1.35	2.94
111812		1.89	3.45		2.53	4.70	3.09	2.93
840894		11.87	1.12	3.23 1.65	2.53 0.60	5.53	5.35	2.92
50114	1	0.28	1.50 3.23	1.03	1.70	5.64	2.05	2.92
971367	9	12.06	1.78	2.25	2.22	3.04	5.31	2.92
1473300 824704		4.26 0.70	3.07	2.49	3.23	2.63	3.15	
82470 ² 897670	1		1	3.98	2.74	2.73	1.43	2.90
43015	1			3.54	2.88	1.69	2.95	2.90
3122	1	1	II .	2.29	4.85	2.20	3.01	2.90
74382	1		· ·	1.72	5.33	1.00	5.28	2.90
152197	•	t	B.	1.94	2.94	3.07	2.92	2.90
51163	1		i	2.68	1.95	4.09	3.56	2.89
36088		•		2.17	4.57	1.22	2.42	2.89
149223			3	3.13	2.20	3.54	2.04	2.89
78483			1	1.72	3.99	1.90	2.77	2.88
76779			B	2.09	2.38	3.81	3.45	2.88
84059		1	4	1.06	2.15	7.48	1.51	2.88
87831		L L	1	2.73	1.60	4.36	3.36	2.87
72548		1		3.28	2.57	3.29	1.57	1
20916		1	1	1.48	3.65	1.17	1.48	2.86
48843	1	1	•	2.58	2.02	1.74	3.84	2.85
73130			2	3.06	2.31	1.95	3.15	2.85
56523	1		1	3.41	1.55	7.11	0.85	1
50429		8 17.89	1.61	2.21	7.54	1.36	1.50	2.84
	-	•	•					

823588	7.72	2.72	2.42	3.62	3.20	2.49	2.44	2.83
878544	19.20	6.78	5.95	1.60	3.38	1.81	1.41	2.83
26736	9.52	3.36	3.45	0.77	0.57	6.03	3.33	2.83
145388	2.66	0.94	3.12	3.26	4.21	2.28	1.26	2.83
76196	2.53	0.90	1.68	2.14	2.06	3.22	5.03	2.83
624271	5.74	2.03	3.08	2.14	1.61	6.02	1.25	2.82
66902	3.21	1.14	3.84	1.85	1.41	0.79	6.22	2.82
86160	2.10	0.74	2.49	3.24	2.74	2.51	3.12	2.82
526945	1.14	0.40	3.22	1.77	2.51	3.16	3.41	2.82
131621	10.21	3.63	3.64	1.77	4.66	1.90	2.11	2.82
815036	1.62	0.58	3.61	3.75	2.37	3.01	1.33	2.81
416833	6.07	2.16	1.37	3.86	1.76	4.96	2.11	2.81
810221	3.85	1.37	1.41	2.35	3.96	2.88	3.43	2.81
1536991	10.79	3.85	4.74	1.63	5.04	1.39	1.21	2.80
1240220	7.60	2.71	0.70	2.40	5.33	4.42	1.16	2.80
125308	2.61	0.93	4.31	2.08	3.64	2.05	1.93	2.80
502690	4.09	1.46	2.11	3.13	2.39	4.17	2.19	2.80
1635874	6.06	2.17	5.71	2.90	1.66	2.16	1.54	2.79
32076	1.03	0.37	1.94	2.97	2.60	3.25	3.21	2.79
856535	1.40	0.50	3.08	3.52	2.81	2.44	2.11	2.79
30793	7.89	2.84	2.22	2.49	3.31	3.04	2.85	2.78
826102	4.19	1.51	1.94	1.78	7.59	1.08	1.50	2.78
356992	21.95	7.93	3.72	2.43	2.23	3.22	2.25	2.77
810571	4.37	1.58	1.62	3.67	1.77	4.25	2.52	2.76
365098	4.93	1.78	4.88	1.79	3.50	1.53	2.10	2.76
788655	7.66	2.78	2.38	1.89	4.08	3.62	1.83	2.76
186918	3.04	1.10	1.49	2.82	2.33	5.76	1.40	2.76
842784	326.54	118.59	0.68	2.49	1.56	5.61	3.44	2.75
127396	14.07	5.12	2.63	3.54	0.99	4.76	1.83	2.75
261472	9.31	3.39	2.65	1.79	2.40	5.00	1.89	2.75
813179	6.60	2.40	1.14	8.50	0.98	2.38	0.73	2.75
240430	3.52	1.28	5.57	1.49	1.68	2.99	1.98	2.74
1435029		0.67	2.78	3.83	3.22	2.31	1.55	2.74
124824	1	11.28	2.73	1.75	1.56	2.43	5.21	2.73
1493390	1	1.31	3.84	1.25	2.88	3.90	1.79	2.73
251936	l .	2.74	3.40	3.01	2.19	2.17	2.89	
343609	1	0.58	2.30	2.63	3.03	2.56	3.13	2.73
125722	1	1.21	4.28	2.15	2.60	3.74	0.81	2.72
322051	5	4.27	2.00	2.69	3.96	1.51	3.38	2.71
756556	1	6.47	1.96	2.33	5.07	2.59	1.60	2.71
430318			1.74	2.76	3.55	1.94	3.54	2.71
795555	L .	1	3.07	3.53	2.44	2.31	2.14	2.70
951125		1	2.18	5.16	1.53	2.32	2.28	2.69
843150		1	2.80	3.47	2.10	1.72	3.38	2.69
5077			1.30	2.66	2.08	5.85	1.54	2.68
8050			0.85	2.49	1.54	4.92	3.61	2.68
13718			1	3.19	2.48	3.67	2.23	2.68
27948		I I		1.33	7.28	0.95	1.69	2.67
74510	1			3.08	3.04	1.84	3.55	2.67
76014			1	2.03	3.48	3.23	2.28	
82361	•	1	1	1.46	5.77	1.58	0.93	
78364			B .	2.53	3.22	1.85	2.18	1
8157		1	3	2.44	4.22	1.59	1.60	1
78579	55.68	3 21.05	3.09	2.27	3.82	1.77	2.28	2.65

arrecal	1416	5.37	1.96	2.26	2.29	3.29	3.39	2.64
277660 66336	14.16	0.63	4.14	4.60	4.12	0.16	1	2.63
1630990	20.64	7.84	4.29	2.18	1.71	4.51	0.48	2.63
247609	3.15	1.20	2.53	1.52	2.16	3.23	3.74	2.63
898123	4.77	1.81	1.92	3.20	3.04	1.92	3.07	2.63
40136	5.23	1.99	1.90	1.77	2.01	1.67	5.77	2.62
46367	1.63	0.62	1.73	1.61	1.24	7.45	1.09	2.62
281072	5.02	1.92	1.08	2.04	1.92	5.46	2.57	2.62
1606837	23.76	9.09	3.80	1.46	1.83	3.18	2.79	2.61
308726	14.73	5.64	3.93	2.00	4.67	0.90	1.56	2.61
784116	0.26	0.10	2.07	3.32	3.40	2.11	2.15	2.61
46620	5.25	2.02	1.48	2.20	1.83	3.69	3.83	2.61
504544	12.31	4.73	1.63	2.83	1.76	5.44	1.35	2.60
279394	3.18	1.23	1.59	1.61	6.58	1.11	2.07	2.59
262865	30.48	11.78	3.45	2.03	4.88	0.98	1.59	2.59
123980	2.38	0.92	1.80	3.18	2.76	1.49	3.71	2.59
293830	4.43	1.72	2.14	1.56	6.45	1.42	1.34	2.58
383188	2.79	1.08	3.03	2.15	1.40	2.92	3.40	2.58
853998	12.39	4.81	1.79	2.00	3.76	2.32	3.01	2.58
813651	4.96	1.93	1.72	2.40	2.48	3.04	3.24	2.58
203350	1.91	0.74	2.76	1.43	3.30	3.24	2.14	2.57
240961	2.11	0.82	2.13	4.67	3.19	1.71	1.17	2.57
1035765	6.21	2.41	2.60	0.83	3.16	0.47	5.78	2.57
970271	3.16	1.23	2.05	3.35	2.83	1.20	3.37	2.56
950836	34.79	13.60	2.31	1.27	6.82	0.77	1.62	2.56
461699	3.44	1.35	1.58	2.10	5.96	1.28	1.84	2.55
743851	•	3.22	3.05	2.50	2.28	3.39	1.54	2.55 2.54
731290	•	4.56	1.16	2.19	1.46	7.52	0.38 2.56	2.52
1475842		89.33	0.55	1.42	1.77	6.27	1.14	2.52
796513			3.08	2.43	2.08	3.84 3.07	3.25	2.50
82905	•		2.88	1.65	1.67	1.14	5.22	2.49
123755	t .	1	3.88	1.17	1.06 6.84	0.91	0.70	2.49
884606			2.43	1.59 1.64	1.66	2.19	3.33	2.48
323611		•	3.60 3.57	1.68	1.35	4.43	1.33	2.47
1472150			3.24	3.20	3.55	1.12	1.25	2.47
950574	•		1	3.06	3.37	2.76	1.82	1
345056 48749		ı	B .	2.00	5.71	2.00	0.89	
28907			l l	2.50	5.68	1.17	1.01	2.45
160434		1	B .	1.51	0.94	6.33	2.34	2.45
89691		1		4.47	0.31	3.53	0.78	2.44
25133			1	1.78	2.05	3.30	3.33	2.44
47011				3.43	1.78	3.57	1.10	1
85467	•		3	7.14	1.43	0.90	1.41	
24586	1	1	3	1.10	4.77	0.61	0.96	1
2843		· ·	1.13	1.64	1.72	5.29	2.3	•
34458	53.2	8 21.92	1.02	3.29	3.08	2.11	2.6	
5119	52 225.6	6 92.93	2 0.39	2.42	0.61	7.65	1.0	
8981		5.9		2.53	3.20	1.41	3.0	
2917	06 2.4		1	1.35	1.24	1.16	6.5	1
2585			B	6.91	0.97		1.8	
14124		t t			0.25		0.2	
14758					1.60		3.2 1.5	
2005	99 13.	42 5.6	3.26	1.35	4.93	0.84	1	2.39

		_			-			
252904	5.40	2.26	3.71	1.41	3.30	1.76		2.39
430614	2.38	1.00	2.34	3.41	1.17	4.02		2.39
381036	3.54	1.49	2.56	1.79	5.52	1.27		2.38
448117	3.91	1.64	3.8 6	1.93	3.01	1.46		2.38
250673	2.66	1.12	5.54	1.03	1.65	1.42		2.37
593929	17.74	7.51	3.61	1.43	5.14	0.93	1	2.36
841615	12.16	5.15	2.59	1.62	5.18	1.30		2.36
52338	2.91	1.24	3.59	1.69	4.60	0.65		2.35
137254	1.35	0.58	3.41	1.79	3.95	1.23		2.35
767994	4.29	1.83	2.20	1.22	7.00	0.55		2.34
197765	8.83	3.78	2.35	0.96	6.16	0.88		2.34
254310	10.26	4.39	2.47	1.99	5.33	1.07		2.33
855244	15.78	6.78	4.53	0.61	5.42	0.42	0.66	2.33
144926	9.20	3.96	3.10	1.65	2.25	3.17	1.43	2.32
565779	31.62	13.68	3.23	1.26	1.97	3.57	1.53	2.31
563451	4.17	1.82	3.24	2.08	3.69	1.02	1.43	2.29
462325	8.76	3.83	1.92	0.60	1.72	5.27	1.93	2.29
132911	52.56	22.96	3.65	1.43	4.04	0.88	1.45	2.29
827168	6.51	2.85	2.17	0.93	6.04	1.22	1.08	2.29
1070015	3.26	1.43	0.69	2.05	0.53	3.30	4.86	2.28
293243	13.17	5.84	3.23	1.48	4.46	0.69	1.43	2.26
743211	3.84	1.71	1.36	1.66	5.64	1.28	1.27	2.24
259462	3.69	1.65	0.66	0.89	0.75	8.24	0.65	2.24
291345	70.40	31.69	5.27	1.32	2.52	0.90	1.10	2.22
281191	7.36	3.32	3.42	1.94	3.36	0.84	1.52	2.22
362080	1	1.72	1.00	0.91	4.67	3.13	1.35	2.21
825058		1.55	2.14	0.93	6.34	1.08	0.54	2.21
290841		2.89	3.60	3.33	1.43	1.62	1.02	2.20
291057		0.61	1.18	1.53	3.04 0.54	4.12 0.73	1.04 8.15	2.18 2.17
855624	3	8.17	0.67 0.80	0.78 0.85	0.34	7.58	0.85	2.17
50344 1056172	1	0.39 4.78	2.40	1.52	0.73	5.66	0.83	2.16
1056203	3	0.89	1.05	0.95	1.43	6.28	1.07	2.15
786609	i i	2.23	0.54	0.55	1.98	7.08	0.62	2.15
248261	1	0.70	3.67	1.05	0.77	0.89	4.36	2.15
42904			1.32	2.03	3.22	1.07	3.04	2.14
74493		I .	0.80	0.91	0.71	7.21	1.06	2.14
46195	ŧ	1	3.10	3.01	1.55	1.32	1.68	2.13
8148	3		1.73	3.05	2.09	0.66	3.02	2.11
89754			6	3.41	2.18	0.69	0.75	2.10
79636	1		1	1.59	3.66	0.93	0.94	2.05
14133	3	B.		0.85	7.18	0.60	0.88	2.05
129165		B .		1.14	3.32	0.94	1.12	2.05
84130	•	3	1	1.26	5.92	0.84	0.91	2.04
62768			1	1.30	5.58	1.43	0.63	2.03
81109	1			1.38	0.73	1.83	0.60	2.02
36194			1	1.71	5.64	0.71	0.95	2.01
7505		1		1.03	1.07	3.28	1.53	2.00
28289			3	1.07	6.06	0.92	0.83	1.99
149290	3.6	8 1.85	1.76	1.20	5.34	0.93	0.72	1.99
2260	0.3	6 0.18	1.38	5.49	0.87	1.20	1.00	1.99
1198:	51 6.5	2 3.28	4.42	3.58	0.65	0.63	0.64	
7842	85 3.9			1.06	3.08	1.24	1.07	1
1407	92 1.6	6 0.8:	5 1.20	1.05	5.74	0.76	1.05	1.96

704076	5.70	2.95	1.28	1.04	3.20	3.25	0.88	1.93
142326	1.49	0.78	5.18	1.78	0.80	0.91	0.92	1.92
840944	79.86	42.51	3.82	0.54	4.16	0.61	0.27	1.88
377672	2.59	1.39	0.54	0.57	0.40	7.29	0.56	1.87
431597	5.45	2.95	1.55	0.82	5.32	0.67	0.90	1.85
32381	2.18	1.18	0.60	0.99	3.26	1.10	3.29	1.85
471641	4.79	2.74	0.55	3.60	0.72	3.36	0.52	1.75
753794	1.65	0.94	0.66	1.32	0.68	5.31	0.78	1.75
950464	9.75	5.67	1.05	0.71	5.37	0.77	0.68	1.72
289832	6.67	4.11	1.15	0.75	5.01	0.74	0.48	1.62
309864		1.62	5.57	0.48	0.62	0.55	0.47	1.54
1492202	1	1.98	0.42	0.35	0.43	0.34	5.02	1.31

Table 1-2

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	ı	GenBank	Aver. of	Aver. of	Aver.	Tumor/	Tumor /	T/N	T/N	T/BPH	T/BPH
	Clone	Acc No.		Normals	of BPH	Normal	BPH	5X, ≥1/5	3X, <u>>2</u> /5	5X, ≥1/5	3X, <u>≥2/5</u>
		AA453562	25.18	0.81	0.50	31.24	50.69	+	+	+.	+
		AA196979	24.42	1.02	0.75	23.91	32.70	+	+	+	+
1		AA779728	33.61	1.51	1.35	22.27	24.97	+	+	+	+
	281003	N50880	128.77	6.30	4.90	20.43	26.26	+	+	+	. +
	308682	W25202	8.99	0.87	0.27	10.37	32.91	+	+	+	+
	281039		14.05	0.94	0.90	14.99	15.57	+	+	+	+
	284701	N64840	33.05	7.05	1.52	4.68	21.68	+	+	+	+
		AA490213	22.39	1.32	2.83	16.99	7.91	+	+	+	+
	42415		6.86	0.51	0.72	13.52	9.49	+	+	+	+
	46166	i	13.13	0.93	1.71	14.12	7.68	+	+	+	+
		AA432103	53.04	7.37	3.83	7.20	13.84	+	+	+	+
		AA427899	12.32	0.65	5.99	18.85	2.05	+	+	-	-
	222157		8.80	1.02	0.72	8.67	12.14	+	+	+	+
		AA043109	24.75	3.47	1.86	7.13	13.32	+	+	+	+
		AA703396	18.37	1.83	1.80	10.05	10.21	+	+	+	+ .
	66336		1.67	0.63	0.10	2.63	16.70	-	+	+	+
	35058	B .	15.59	1.89	1.44	8.24	10.81	+	+	+	+
	26736	l .	9.52	3.36	0.62	2.83	15.37	+-	+	+	+
		AA159714	28.06	5.61	2.29	5.00	12.23	+	+	+	+
	32160	li .	26.80	3.59	2.77	7.47	9.69	+	+	+	+
orar :	814053	AA465495	14.66	1.70	1.75	8.61	8.38	+	+	+	+
100 100 100 100 100 100 100 100 100 100	292515	4	16.22	2.86	1.56	5.67	10.38	+	+	+	+
		AA436097	36.49	7.10	3.45	5.14	10.57	+	+	+	+
	282089		11.18	1.51	1.41	7.41	7.92	+	+	+	+
	283313	N54763	63.94	6.63	12.14	9.65	5.27	' +	+	+	+
2.2.4	29125	L .	15.20	1.35	4.65	11.29	3.27	/	+	+	+
	104881	0 AA621342	20.15	3.04	2.59	6.63	7.78	3 +	+	+	+
222	19752	5 H52119	9.52	1.85	1.06	5.16	9.01	+	+	+	+
	6740	1 T49326	1.47	0.20	0.23	7.28	6.27	7 +	•	+	-
1784 1874 1876	49017	8 AA121266	24.55	5.63	2.73	4.36	8.99	9 +	-	+	+
Contraction of the contraction o	155652	6 AA935560	0.88	0.15	0.12	5.96	5 7.33	3 +	•	+	-
S H	74615	2 AA419486	9.35	2.23	1.07	4.18			+	+	+
	103473	8 AA780190	22.40					1	+	+	+
2000	25937	14 N31952	10.51	2.33					+	+	+
To the second se		35 AA437224							+	+	. +
Fails	75428	80 AA479284							+	+	+
		35 AA625634							+	+	+
	7129	50 AA28227:							+	+	+
	507	72 H1680							-	+	+
		08 AI00027						•	-	+	+
		94 AA67941							+	-	+
		64 AA04726						1	•	+	+
		81 AA84515							-	+	-
	2831							L	+	+	+
		67 AA45496						30 +	+	-	
	265							59 -	-	+	+
		297 AA23522						48 -	+	+	
		34 AA23224					07 10.		-	+	+
		260 AA88492						96 +	+	+	+
	2583							.86 +	+	+	+
		279 AA6829						45 +	+	- +	- +
	1610	448 AA9918	56 135.	55 57.0	06 16.4	14 2.	38 8.	.25 -	-	т	7

Page 1

461522	A705237	1.21	0.65	0.14	1.87	8.74	-	-	+	+
	AA633957	8.99	2.01	1.47	4.48	6.12	+	+	+	+
280217	N47941	20.52	5.86	2.89	3.50	7.09	+	+	+	+
240694	H78135	4.53	0.90	0.83	5.01	5.47	+	+	+	+
321706	W35398	7.80	1.86	1.25	4.20	6.23	+	+	+	+
1	AA430545	11.43	2.60	1.93	4.40	5.94	+	+	+	+
	AI732784	19.62	2.74	6.22	7.16	3.15	+	+	-	+
51773	H23211	11.48	2.40	2.09	4.77	5.50	+	+	+	+
66317	T66815	5.50	1.76	0.78	3.13	7.06	+	+	+	+
35626	R45292	9.42	1.09	6.51	8.67	1.45	+	+	-	-
	AA682851	9.39	1.62	2.25	5.80	4.17	+	+	+	+
	AA702781	14.83	7.21	1.88	2.06	7.90	•	-	+	+
1	AA416843	11.60	4.56	1.57	2.54	7.40	+	•	+	+
	AA504783	4.27	2.51	0.52	1.70	8.20	•	-	+	+
	AA136125	35.83	12.59	5.15	2.85	6.96	+	+	+	+
	AA481464	47.06	7.13	14.79	6.60	3.18	+	+	•	+
293635	N63807	26.47	3.44	12.94	7.69	2.05	+	+	-	-
757392	AA426110	0.86	0.37	0.12	2.32	7.41	•	-	+	+
	AA173189	3.58	2.33	0.44	1.54	8.16	-	-	+	+
151055	H02230	27.32	7.51	4.62	3.64	5.92	+	+	+	+
297439	W03687	48.60	11.33	9.26	4.29	5.25	+	+	+	+
1	AA457172	11.15	1.82	3.27	6.12	3.41	+	+	-	+
	AA677629	0.87	0.46	0.11	1.87	7.58	-	•	+	+
	AA173598	3.07	2.45	0.38	1.25	8.17	-	•	+	+
	AI025015	5.68	0.93	1.77	6.13	3.21	+	+	-	+
	AA129551	12.35	3.96	1.99	3.12	6.21	+	•	+	+
293651	1	13.35	1.84	6.56	7.26	2.04	+	+	-	-
	AA625666	2.15	0.29	1.14	7.39	1.88	+	+	-	-
	AA404486	113.14	18.10	38.24	6.25	2.96	+	+	-	+
262053	1	10.75	5.15	1.52	2.09	7.08	-	-	+	+
81331	i	2.84	0.78	0.51	3.63	5.53	+	-	+	+
840894	AA482340	34.81	11.87	5.66	2.93	6.15	-	+	+	+
839081	AA488732	16.60	5.37	2.78	3.09	5.98	+	+	+	+
41790	W90519	12.10	2.57	2.82	4.71	4.29	+	+	+	+
796366	6 AA459761	34.94	17.03	5.05	2.05	6.92	•	+	+	+
35966	1 AA011095	26.24	8.79	4.39	2.99	5.97	-	+	+	+
5018	8 H17943	10.75	2.29	2.53	4.69	4.25	+	+	-	+
8495	5 T74846	5.26	1.01	1.43	5.21	3.69	+	+	+	+
15960	8 H16152	14.43	4.14	2.69	3.49	5.36	+	+	+	+
139058	4 AA843592	6.54	1.91	1.21	3.42	5.41	+	+	+	+
82573	6 AA504842	0.92	0.60	0.13	1.54	7.26	-	•	+	+
2216	1 T66154	20.24	13.40	2.80	1.51	7.24	-	•	+	+
78793	8 AA452278	5.09	1.27	1.08	4.01	4.71	+	+	+	+
80960	3 AA442991	49.03	6.94	29.52	7.06	1.66		+	-	-
82610	9 AA521327	0.75	0.54	0.10	1.38	7.27	1	•	+	+
88460	06 AA630006	33.46	13.42	5.43	2.49	6.16	3	-	+	+
82582	22 AA491370	14.07	9.48	1.97	1.48	7.16		-	+	+
	28 AA070495		3.89	6.62	5.44	3.20		+	-	+
8140	17 AA455668		23.42	17.57	3.68	4.90		+	+	+
8140	62 AA465338	0.81	0.47	0.12	1.72	6.83	1	-	+	+
	03 AA521416		0.82	0.19	1.61	6.93		•	+	+
	70 AA779919		1.72	2.71	5.19	3.28	1	+	•	+
4157			0.61	1.15	5.53	2.93		+	•	+
8151	63 AA481144	1.20	0.57	0.19	2.09	6.30	5 -	-	+	+

					4.00	1				
	AA464568	14.49	3.00	4.01	4.83	3.61	+	+	+	+
	AA464689	2.17	0.39	0.77	5.60	2.82	+	+	•	+
	AA010932	7.82	2.50	1.49	3.13	5.26	+	+	+	+
	AA451807	4.58	1.26	0.97	3.63	4.74	+	-	+	+
840306	AA485333	2.49	1.26	0.39	1.97	6.38	-	-	+	+
212165	H68845	41.96	8.42	12.58	4.99	3.34	+	+	+	+
138444	R68272	4.04	1.00	0.95	4.06	4.24	+	+	+	+
824723	AA488986	14.11	4.37	2.84	3.23	4.96	+	+	+	+
186918	H43317	3.04	1.10	0.56	2.76	5.38	+	-	+	+
191904	H38880	7.67	1.44	2.75	5.32	2.79	+	+	-	+
		20.64	7.84	3.79	2.63	5.44	-	+	+	+
	AA608567	52.65	11.99	14.32	4.39	3.68	+	+	+	+
	AA476234	9.44	1.78	3.46	5.31	2.73	+	+	_	-
	AA521394	1.20	0.83	0.18	1.44	6.59	_	-	+	+
30428	R42061	4.29	0.99	1.16	4.35	3.68	+	+	+	+
	AA620995	10.33	4.78	1.77	2.16	5.84	+	_	+	+
251806				0.46	1.60	6.39		_	+	+
	H96647	2.94	1.83				-	+	+	+
305538	N89861	11.44	3.31	2.54	3.46	4.50	-	т	+	+
52933	H29315	3.59	1.99	0.58	1.80	6.14	-	-	+	+
587415	AI733857	12.98	3.44	3.12	3.77	4.16	+	+		
	AA490522	1.38	1.09	0.21	1.27	6.65	•	<u>-</u>	+	+
416436		11.08	2.37	3.42	4.67	3.24	+	+	+	+
321470		7.74	1.68	2.36	4.61	3.29	+	+	•	+
842925	AA486444	4.61	1.21	1.13	3.79	4.07	-	+	-	+
859228	AA666366	85.06	37.74	15.20	2.25	5.60	•	-	+	+
845363	AA644092	29.59	7.07	8.08	4.18	3.66	+	+	+	+
362755	AA018675	16.82	3.11	6.91	5.41	2.43	+	+	-	-
253132	H89036	1.40	1.21	0.21	1.16	6.66	-	-	+	+
233419	5	5.55	1.14	1.90	4.87	2.93	+	+	-	+
	AA953229	5.30	1.51	1.24	3.52	4.28	-	+	-	+
	AA608636	3.00	1.87	0.48	1.60	6.18	-	-	+	+
	AA620591	6.84	6.68	1.01	1.03	6.75	-	-	+	+
358850	1	21.29	5.98	5.06	3.56	4.21	+	+	+	+
	8 AA427395	10.54	2.90	2.58	3.63	4.08	_	+	+	+
	4 AA486533	79.86	42.51	13.69	1.88	5.83		+	+	+
26821		3.43	0.70	1.23	4.93	2.78	+	+	+	-
	6 AA487914	15.85	7.18	2.88	2.21	5.50	_	_	+	+
	1 AA461092	30.31	5.27	15.52	5.75	1.95	+	+	-	_
26610	•	20.68	4.53	6.63	4.57	3.12	+	+	+	+
	T .		5.34	1.75	1.90	5.78	:		+	+
	2 AA458516	10.13	16.50	17.00	3.90	3.78	+	+	+	+
	7 AA402352	64.30			3.90	4.66	1	+	+	+
21400	II .	4.67	1.55	1.00				+		_
	2 AA777488	42.54	7.72	19.71	5.51	2.16	1	+	-	+
15026		13.66	3.31	3.87	4.12	3.53	1		-	
28989		7.82	1.84	2.30	4.25	3.40	1	+	+	+
	31 AA047441	10.48	3.68	2.19	2.85	4.78		+	+	+
	55 AA449837	7.66	2.78	1.58	2.76	4.86		+	+	+
20808	1		1.98	2.19	3.99	3.60	3	+	+	+
	03 AA625995		11.14	11.32	3.83	3.77	8	+	+	+
	31 AA045699		23.94		1.65	5.95	1	-	+	+
	63 AA 136386		2.97		1.32	6.27	E .	-	+	+
	87 AA054701		7.39		5.34	2.25		+	-	-
	46 AA461467		4.35		1.43	6.16		-	+	+
8142	35 AA465598	12.37	6.41	2.19	1.93	5.65	5 -	-	+	+
	-									

				•	CDIO I O	_				
53022	R15880	1.34	0.70	0.24	1.90	5.63	•	•	+	+
377246	AA055306	12.60	3.02	3.78	4.17	3.34	+	+	+	-
814662	AA481045	1.12	0.67	0.19	1.67	5.84	-	-	+	+
45327	H08548	8.18	2.30	2.08	3.56	3.94	•	+	+	+
51700	H22853	3.66	1.21	0.82	3.02	4.47	-	+	+	+
80050	T63893	8.15	3.84	1.52	2.12	5.35	-	•	+	+
	AA172372	14.61	10.98	2.38	1.33	6.14	-	•	+	+
795907	AA460346	5.39	1.25	1.71	4.31	3.15	+	+	+	+
824126	AA490611	0.59	0.37	0.10	1.58	5.87	-	•	+	+
	AA100674	225.66	92.92	45.17	2.43	5.00	+	-	+	+
	AA496957	9.40	3.54	1.97	2.65	4.76	+	+	+	+
	AA432075	17.23	6.93	3.50	2.49	4.93	-	-	+	+
	AA918370	401.67	285.67	66.91	1.41	6.00	•	-	+	+
	AA988746	9.00	4.61	1.65	1.95	5.45	-	-	+	+
	AA996028	23.76	9.09	4.98	2.61	4.77	-	+	+	+
	AA495819	14.29	3.54	4.28	4.03	3.34	+	+	-	+
796239	AA460675	26.07	23.22	4.18	1.12	6.24	-	-	+	+
49302	1	4.88	0.86	2.90	5.68	1.68	+	+	-	-
	AA436142	12.77	7.46	2.26	1.71	5.64	-	-	+	+
207029	3	81.88	14.79	45.41	5.54	1.80	+	+	-	-
	AA458747	1.34	1.36	0.21	0.99	6.35	-	•	+	+
1636108		3.58	1.45	0.74	2.47	4.85	-	-	+	+
	4 AA460719	0.66	0.83	0.10	0.80	6.50	-	-	+	+
23882	1	6.54	2.22	1.50	2.94	4.35	+	+	+	+
12902	1	4.30	1.04	1.36	4.13	3.16	+	+	+	+
	3 AA430718	16.14	6.63	3.34	2.43	4.83	-	-	+	+
28907	1	37.48	15.32	7.78	2.45	4.82	+	-	+	+
163728	li .	4.01	2.58	0.70	1.55	5.71	-	-	+	+
26369		4.23	0.93	1.58	4.57	2.68	+	+	-	+
	7 AA683050	35.22	12.06	8.15	2.92	4.32	+	+	+	+
3238	1	2.18	1.18	0.41	1.85	5.38	-	+	+	+
	30 AA130677	1.92	1.75	0.31	1.10	6.12	-	-	+	+
	36 AA465148		0.58	0.37	2.81	4.40	-	+	+	+
	85 AA074258		3.46	3.22	3.48	3.73	+	+	+	+
4163	9		2.66	0.81	1.69	5.52	-	-	+	+
	66 AA448402		3.84	2.97	3.14	4.05	+	+	+	+
2136			4.10	3.27	3.19	3.99	+	+	+	+
	07 AA884713		0.99	1.07	3.72	3.44	-	+	-	+
	82 AA032221		12.66	6.60	2.45	4.70	-	-	+	+
	76 AA055880		56.58	27.52	2.34	4.81	-	•	+	+
2429			4.39	5.16	3.86	3.28	-	+	-	+
	27 AA775422		0.87	1.16	4.07	3.05	+	+	-	+
	68 AI79117			4.21	4.23	2.87	+	+	+	+
2877				0.89	3.97	3.10	+	+	+	+
	927 AA48644			0.72	1.18	5.88	-	-	+	+
	042 AA67676			1.45	3.63	3.42	+	+	+	+
	281 AA67028			1.37	2.06	4.97	-	-	+	+
	701 AA01117			1.09	2.25	4.78	3 -	-	+	+
	750 N3623			5.70	3.40	3.61	l	+	+	+
	819 AA49039			3.34	3.41	3.58	3 -	+	-	+
	807 AA59853			0.98	1.83	5.10	5 -	-	+	+
	749 AA66398				4.20	2.7	7 +	+	-	-
	895 N2117				4.82			+	-	-
	207 AA50412			4.19	4.09	2.8	6	+	-	-
230							-			

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461476	A705041	0.78	0.61	0.14	1.28	5.64	-	-	+	+
323474	W45623	1.25	0.32	0.41	3.87	3.06	+	+	+	+
744846	AA625675	2.69	0.70	0.87	3.83	3.10	+	+	•	+
741474	AA401111	4.93	1.40	1.45	3.51	3.40	+	+	+	+
506564	AA709044	12.93	4.03	3.49	3.21	3.70	-	+	+	+
449275	AA777705	3.81	0.83	1.64	4.58	2.32	+	+	+	-
878544	AA775863	19.20	6.78	4.73	2.83	4.06	+	+	+	+
1	AA422139	17.93	9.29	3.62	1.93	4.95	<u>-</u>	-	+	+
259973	N47829	1.66	1.95	0.28	0.85	5.99	-	-	+	+
290370	N62301	0.62	0.48	0.11	1.30	5.53	-	-	+	+
	AA425534	15.94	8.17	3.27	1.95	4.87	•	-	+	+
1	AA406505	7.78	2.14	2.45	3.63	3.18	-	+	•	+
290841	N99711	6.35	2.89	1.38	2.20	4.61	-	+	+	+
50606	H17763	14.94	6.00	3.47	2.49	4.31	-	•	+	+
	AA873159	5.15	1.21	2.03	4.26	2.53	+	+	+	-
252314	H87175	0.69	0.62	0.12	1.11	5.66	-	-	+	+
	AA455940	1.75	0.86	0.37	2.04	4.73	-	-	+	+
	AA620556	18.86	7.00	4.63	2.69	4.07	+	-	+	+
	AA677165	20.56	33.95	3.34	0.61	6.16	-	-	+	+
	AA699427	7.29	2.43	1.95	3.00	3.74	+	+	+	+
	AA976714	6.44	3.55	1.31	1.81	4.92	-	-	+	+
	AA150502	0.81	0.59	0.15	1.36	5.37	-	-	+	+
	AA133044	559.22	179.15	155.61	3.12	3.59	+	+	+	+
245547	L .	5.86	1.63	1.88	3.59	3.12	+	+	+	+
200599	1	13.42	5.61	3.11	2.39	4.32	-	+	+	+
139771	•	7.72	6.17	1.42	1.25	5.44	-	-	+	+
	AA705376	8.32	6.33	1.55	1.31	5.38	-	-	+	+
	AA464580	4.37	1.58	1.11	2.76	3.92	-	+	+	+
279482	I .	8.66	3.25	2.16	2.67	4.00	+	-	+	+
127396	1	14.07	5.12	3.61	2.75	3.90	-	+	+	+
	AA490609	4.64	2.75	0.94	1.69	4.95	-	-	+	+
	3 AA609323	38.01	9.06	15.59	4.20	2.44	+	+	+	-
	AA287375	9.94	3.00	3.00	3.31	3.31	-	+	•	+
25086		123.92	32.54	44.10	3.81	2.81	+	-	+	-
163174		5.09	1.33	1.84	3.84	2.77	+	+	•	+
	4 AA779391	18.03	7.40	4.35	2.44	4.14	-	+	+	+
	2 AA609368	7.90	4.94	1.59	1.60	4.98	-	-	+	+
8157	4	12.77	4.82	3.25	2.65	3.93	-	+	+	+
	4 AA434360	103.13	30.63	32.18	3.37	3.20	+	+	+	+
	5 AA621761	14.86	3.64	5.98	4.08	2.48	+	+	-	-
	6 AA459100	18.28	10.68	3.77	1.71	4.85	-	-	+	+
7791		14.99	17.48	2.63	0.86	5.70	-	-	+	+
26011		0.88	0.86	0.16	1.03	5.52	-	-	+	+
	53 AA458551	17.84	5.98	5.02	2.98	3.55	-	+	-	+
21040		34.02	7.84	15.49	4.34	2.20	+	+	-	-
12482	1	30.85	11.28	8.13	2.73	3.79	+	-	+	+
	29 AA169379	17.74	7.51	4.26	2.36	4.17	+	+	+	+
	43 AA278473		2.81	0.99	1.70	4.82	-	-	+	+
	33 AA058818		6.81	3.48	2.20	4.32	2 -	-	+	+
1916	1		4.64		4.77	1.74	+	+	-	-
	85 AA012866		13.93		2.89	3.6	1 -	+	+	+
	67 AA491265		0.41		1.81	4.6	1	•	+	4
	71 AA181179				2.82			+	+	4
	00 AA885642		3.34		1.83		3 -	•	+	4
-5000	- 1						-			

813973 A	A455689	3.24	0.88	1.18	3.70	2.75	-	+	•	•
	A504536	0.96	0.65	0.19	1.48	4.97	•	•	+	+
592403 A	A159497	0.46	0.16	0.13	2.95	3.50	+	-	+	+
838807 A	A457681	1.90	2.58	0.33	0.74	5.71	•	-	+	+
845352 A	A644080	13.36	5.39	3.37	2.48	3.97	-	-	+	+
814798 A	AA465614	32.86	15.63	7.58	2.10	4.34	-	-	+	+
1607765 A	A989473	3.96	1.73	0.96	2.29	4.14	-	-	+	+
1035765 A		6.21	2.41	1.61	2.57	3.86	+	+	+	+
824739 A	AA488784	2.63	1.02	0.68	2.57	3.86	-	-	+	+
290642	N99659	7.76	2.56	2.28	3.02	3.40	-	+	•	+
121489	T97303	34.70	10.55	11.09	3.29	3.13	-	+	•	+
•	AA489647	7.42	4.65	1.54	1.59	4.82	-	-	+	+
86160	T72336	2.10	0.74	0.59	2.82	3.58	-	+	•	+
	AA862465	41.61	40.32	7.76	1.03	5.36	-	-	+	+
	AA598470	2.07	0.49	0.98	4.25	2.12	+	+	-	•
	AA488238	67.93	15.10	36.21	4.50	1.88	+	+	-	-
1	AA149096	12.31	4.73	3.28	2.60	3.76	+	•	+	+
	AA883679	1.21	0.71	0.26	1.71	4.63	-	-	+	+
1	AA598483	6.77	3.15	1.62	2.15	4.19	-	-	+	+
280763	N50563	0.53	0.50	0.10	1.05	5.28	-	-	+	+
1	AA875936	13.96	4.83	4.08	2.89	3.43	-	+	•	+
194364	H50770	5.44	1.19	3.10	4.56	1.76	+	+	-	-
I I	AA699895	3.24	0.96	1.10	3.36	2.96	-	+	-	+
	AA430351	41.57	23.92	9.10	1.74	4.57	•	.•	+	+
29063	R40970	3.34	1.66	0.78	2.01	4.29	-	•	+	+
	AA171459	1.70	1.73	0.32	0.98	5.32	-	-	+	+
26617	R39862	4.62	2.68	1.01	1.73	4.56	-	-	+	+
	AA045115	4.41	1.79	1.16	2.47	3.82	+	•	+	+
1631849	AI004315	25.01	12.24	5.91	2.04	4.24	-	-	+	+
204111	H55907	3.38	0.87	1.40	3.86	2.42	+	+	- +	-
	AA432312	9.75	3.24	2.99	3.01	3.26 2.80	++	+	+	+
	AA137144 AA486145	6.35 23.03	1.83 5.46	2.27 11.26	3.47 4.22	2.05	+	+	_	τ -
	AA082943	23.03 9.57	3.46 4. 0 0	2.48	2.39	3.86	_	•	_	+
	AA446906	9.37 11.76	4.00 3.47	4.12	3.39	2.85	-	+	_	+
196189	1	11.70	4.82	2.86	2.32	3.91	-	_	_	+
47542	E .	8.07	1.88	4.18	4.30	1.93	+	+	_	_
47384	1	4.22	1.17	1.61	3.61	2.62		+	_	-
	AA491249	74.89	31.45	19.54	2.38	3.83	_		+	+
194723	1	5.22	3.86	1.07	1.35	4.86]	_	+	+
150314		5.29	2.31	1.35	2.29	3.92	_	-	+	+
	AA486200	326.54	118.59	94.88	2.75	3.44	+	+	+	+
48286		2.34	0.61	0.99	3.83	2.36		+	-	-
	1 AA677287	2.69	0.70	1.13	3.82	2.38	+	+	-	-
2420	1	7.49	3.51	1.85	2.13	4.05	-	-	+	+
27354	1	9.10	3.09	2.81	2.95	3.23		+	-	+
26741	1	0.97	0.96	0.19	1.01	5.16	-	-	+	+
	0 AA412250	0.53	0.58	0.10	0.92	5.24		•	+	+
	3 AA977296	5.17	1.57	1.81	3.30	2.85	1	+	-	+
	8 AA505063	2.26	0.96	0.60	2.36	3.77		-	-	+
26286		36.97	16.43	9.52	2.25	3.88	3	-	+	+
	1 AA626146	30.25	19.90	6.55	1.52	4.62	•	•	+	+
86945	0 AA680244	51.58	35.87	10.98	1.44	4.70	-	•	+	+
3798	0 R61372	0.84	0.29	0.27	2.96	3.18	; +	•	+	-

Table 1-3

						_				
124252	R02439	6.06	4.74	1.25	1.28	4.85	-		+	+
238661	H81543	2.74	2.18	0.56	1.26	4.86	•	-	+	+
593658 A		1.08	1.02	0.21	1.07	5.04	-	-	+	T
785616 A		11.47	3.12	4.70	3.67	2.44	-	+	- +	+
84022	T70901	6.97	2.71	1.97	2.57	3.54	-	- +	т	+
511632 A		7.85	2.71	2.46	2.89	3.20	•	T	- +	+
34134	R44734	2.73	1.33	0.68	2.06	4.03	•	+	-	+
39885	R52543	0.94	0.30	0.31	3.11	2.98 3.20	- +	-	+	
840590 A		21.70	7.54	6.78	2.88	3.89	7	-	-	+
49710	H28973	3.21	1.47	0.82	2.18 3.66	2.41	+	+	+	•
42803	R60014	2.96	0.81	1.23	3.00 1.48	4.58	_	•	+	+
134969	R31789	6.81	4.59	1.48 13.09	2.11	3.95	_	_	-	+
75415	T57609	51.71	24.46		3.78	2.28	_	+	-	-
843094 A		49.18	13.00	21.59	2.68	3.37	_	+	_	+
137189	R36409	5.26	1.96	1.56	3.10	2.94	_	+		+
45578	H08016	5.05	1.63	1.72 2.30	3.62	2.41	+		+	•
	A680186	5.55	1.53	4.50	2.53	3.51	_	_	+	+
67654	T49539	15.79	6.24	30.93	3.49	2.53	+	+	_	-
	AA102591	78.20	22.40 1.80	0.50	1.30	4.72		· -	+	+
	AA459765	2.34		6.23	1.92	4.09	_	-	+	+
1637296		25.52	13.27	17.02	5.26	0.75	+	+	-	-
	AA083485	12.81	2.43	0.11	1.42	4.58	_		+	+
	AA489696	0.49	0.34	4.63	1.42	4.61	_	_	+	+
3	AA425692	21.34	15.28 2.23	1.03	1.89	4.10	_	-	+	+
	AA679569	4.23		2.50	2.16	3.84	-	_	+	+
	AA481256	9.59	4.44 6.59	2.52	1.66	4.33	-	-	+	+
196650	R93237	10.93 26.71	9.34	8.54	2.86	3.13	-	+	-	+
	AA398521		0.83	1.23	3.58	2.40	+	+	-	-
	AA872436	2.96 6.36	4.30	1.23	1.48	4.50	_	-	+	+
1	AA485748	6.22	1.68	2.75	3.72	2.26	-	+	-	-
	AA062805	16.03	15.09	3.27	1.06	4.91	-	-	+	+
	AA287067 AA442301	10.03	6.52	3.19	1.96	4.01	_	-	-	+
	AA788970	7.60	2.71	2.40	2.80	3.17	+	+	+	+
210862	i	2.56	1.16	0.68	2.21	3.76		-	+	+
	AA083671	8.46	7.83	1.73	1.08	4.88	-	-	+	+
	AA425773	21.90	19.94	4.51	1.10	4.86	_	-	+	+
	AA405597	1.03	1.35	0.20	0.76	5.19	-	-	+	+
	AA460251	17.83	7.09	5.20	2.51	3.43	-	+	+	+
66902	li .	3.21	1.14	1.03	2.82	3.12	+	+ +	+	+
	AA505162	4.03	1.32	1.40	3.06	2.88	-	+	-	+
201282		6.42	4.16	1.46	1.55	4.39	-	-	+	+
	AA987644	1.97	0.81	0.57	2.45	3.48		-	+	+
	AA018979	4.39	2.04	1.16	2.15	3.77	-	-	+	+
	1 AA864875	12.82	5.49	3.58	2.33	3.58	-	-	-	+
	SAA599717	3.26	1.43	0.90	2.28	3.63	-	+	+	+
10915	R .	5.46	1.70	2.03	3.20	2.69	-	+	-	•
5076		0.43	0.23	0.11	1.89	4.00) -	-	+	+
35699	1		7.93	7.04	2.77	3.12	2 -	+	•	+
	4 AA996104		2.17	1.97	2.79	3.07	7 +	-	+	+
	3 AA488865		2.24		1.80	4.06	5 -	-	+	+
25946			1.65		2.24	3.62	1	-	+	-
	8 AA425205		17.78		1.49	4.37		-	+	+
	O AA 190850		3.82	1.89	1.93	3.9	1 -	-	+	+
	•									

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	743749	AA634291	5.10	1.65	1.86	3.10	2.74	-	+	-	-
	79000	T62164	6.24	3.53	1.54	1.77	4.05	-	-	+	+
	23576		3.58	1.59	1.00	2.25	3.57	-	-	•	+
	366815	AA029490	71.61	18.49	36.97	3.87	1.94	+	-	+	-
	731016	AA421256	0.85	1.00	0.17	0.85	4.96	-	-	+	+
	415437		3.52	1.02	1.50	3.46	2.35	-	+	•	•
	197779	i e	8.61	6.61	1.91	1.30	4.50	-	-	+	+
		AA902831	0.85	0.55	0.20	1.55	4.25	-	•	+	+
		AA126009	24.31	24.09	5.08	1.01	4.79	-	-	+	+
		AA181995	6.17	2.49	1.86	2.47	3.32	-	-	•	+
		AA456616	24.75	12.84	6.40	1.93	3.87	-	-	+	+
		AA015658	28.16	15.79	7.03	1.78	4.01	-	•	+	+
		AA682627	13.81	4.59	4.98	3.01	2.77	_	+	-	+
		AA280288	0.50	0.30	0.12	1.68	4.10	-	-	+	+
	296788	•	3.27	0.67	3.50	4.85	0.93	+	+	•	-
	1638852		1.21	0.50	0.36	2.41	3.36	-	•	-	+
	1636868		5.07	2.89	1.26	1.76	4.01	-	•	+	+
		AA705142	11.87	12.89	2.45	0.92	4.84	-	-	+	+
	357364		0.72	0.27	0.23	2.67	3.09	_		+	+
		AA169645	24.81	12.50	6.57	1.98	3.78	_	_	+	+
		AA486412	2.20	1.38	0.53	1.60	4.16	_	_	+	+
	111812		5.56	1.89	1.98	2.94	2.81	_	+		+
ŧ		AA443969	5.25	4.09	1.17	1.28	4.47	_	,	+	+
ě	292042	1	9.39	2.67	4.19	3.51	2.24	-	+		
ë E		8 AA634379	3.34	1.15	1.17	2.90	2.85	+	+	+	+
e di		7 AA629910	6.68	2.72	2.04	2.46	3.28		<u>.</u>	_	+
# #		4 AA634360	5.09	1.90	1.66	2.68	3.06	-	-	_	+
) 1		9 AA019445	3.03	1.24	0.92	2.44	3.30	_	<u>-</u>	+	+
1		0 AA127100	4.09	1.24	1.39	2.44	2.93	-	+	4	+
1		8 AA630084	6.21	2.55	1.89	2.44	3.29	+	т	+	
152		l l	15.94	8.27	4.20	1.93	3.79	т	-	+	+
_		5 AA189113		1.53		2.72	3.00	•	-	+	+
1111		2 AA021586	4.15		1.39		4.79	•	-	+	+
=	26446	•	0.77	0.84	0.16	0.91		-	-	Ŧ	+
381181	23262	•	4.02	1.27	1.58	3.16	2.55 2.14	+	+	-	7
=		0 AA599064	3.18 2.59	0.89 2.11	1.49 0.58	3.56 1.23	4.47	+	•	+	+
		51 AA457767 52 AA400732				1.60	4.47	-	-		+
Ē		t .	2.81	1.76	0.69		2.62	-	+	+	+
	25133	11 AA169192	34.63	11.27 3.78	13.23	3.07	3.25	-	+	-	+
			9.21	2.72	2.83 2.70	2.44 2.83		•	+	•	+
		88 AA497127	7.72				2.85	-	т	+	+
		48 AA017383	3.13	1.69	0.82	1.85	3.83		-		т
		02 AA521409	4.19	1.51	1.45	2.78	2.89	+	-	+	-
		32 AA625628	44.43	13.65	18.48	3.25	2.40	1 :	+	•	•
		47 AA630800		2.09	6.13	4.22	1.44	+	+	-	-
	337	L .		0.37	0.71	3.73	1.93	-	+	-	+
		49 AA456827		1.97	0.51	1.17	4.49	-	-	+	+
		43 AA495936		15.35	6.79	1.73	3.92	-	-	+	
		02 AA864681		2.54	1.88	2.40	3.25	1	<u>-</u>	-	+
		03 AA626365		3.33	2.98	2.67	2.97	1 .	+	-	+
		58 AA447018		2.77	5.04	3.64	2.00	+	+	-	•
		36 AA608679		13.60		2.56	3.07	B .	-	+	-
		52 AA176607		0.87		0.99	4.63		•	+	+
		143 AI015359		4.49		1.12	4.50	4	-	+	+
	4023	325 AA705516	8.76	3.83	2.63	2.29	3.33	+	-	+	•

Fable	1-3

878284	AA670280	4.41	2.26	1.20	1.95	3.66	-	-	+	+
	AA042812	7.64	5.06	1.86	1.51	4.10	•	-	+	+
	AA670296	5.34	1.86	1.95	2.87	2.73	-	+	-	+
251019	H97778	45.62	22.03	12.95	2.07	3.52	-	-	+	+
258167	N40582	17.24	8.15	4.96	2.12	3.48	•	-	+	+
416833	W87312	6.07	2.16	2.18	2.81	2.78	-	+	-	+
186768	H51914	9.43	5.75	2.39	1.64	3.95	-	-	+	-
1	AA976063	8.20	5.56	1.99	1.47	4.11	•	-	+	+
49728	H29198	1.27	0.57	0.38	2.22	3.36	-	-	-	+
125722	R07506	3.28	1.21	1.14	2.72	2.87	-	+	-	+
	AA181767	1.07	0.78	0.25	1.37	4.22	-	-	+	+
	AA112515	1.14	0.40	0.41	2.82	2.77	•	+	-	+
79032	3	37.58	31.85	8.53	1.18	4.40	-	-	+	+
	AA485922	39.96	18.13	11.84	2.20	3.37	-	-	-	+
	AA487213	7.41	3.64	2.09	2.04	3.54	+	-	+	-
191978		47.95	29.72	12.11	1.61	3.96	-	-	+	+
135900	I B	9.44	5.00	2.57	1.89	3.68	-	-	+	+
	AA282983	2.44	0.98	0.79	2.48	3.08	-	-	-	+
	AA191512	8.33	4.14	2.35	2.01	3.54	•	-	+	+
322914	li .	9.34	3.68	3.10	2.54	3.02	-	•	•	+
251517	1	0.73	0.80	0.16	0.91	4.64	-	-	+	+
	AA482228	25.81	27.85	5.59	0.93	4.61	•	-	+	+
	AA191245	21.60	13.98	5.41	1.55	3.99	-	-	+	+
291345	l .	70.40	31.69	21.23	2.22	3.32	+	-	+	+
80500	I .	140.81	52.48	49.33	2.68	2.85	-	+	+	+
	7 AA485201	3.64	2.20	0.94	1.65	3.88	_	-	+	+
	9 AA703079	168.72	131.76	39.68	1.28	4.25		-	+	+
	9 AA705077	0.62	0.38	0.16	1.66	3.87	_	_	+	+
		4.05	3.19	0.10	1.27	4.25	<u> </u>	-	+	+
36.7	8 AA001234 8 R37224	5.06	1.71	1.97	2.95	2.57	_	+	-	+
13763	4 AA664101	3.00 17.75	8.17	5.31	2.17	3.34	+	-	+	-
	3	12.35	4.01	5.08	3.08	2.43	+	+	+	•
10931	6 T80924 8 AA399198	4.92	2.96	1.28	1.66	3.85		-	+	+
	8 AA708342	34.95	10.61	15.79	3.29	2.21	١.	+	-	-
		19.60	12.65	4.95	1.55	3.96	_	-	+	+
12070	1	1.46	1.23	0.34	1.18	4.32	4	-	-	+
	20 AA486723	21.83	14.39	5.47	1.52	3.99	1	-	+	+
	94 AA994757 98 AA418755	5.70	1.98	2.18	2.88	2.61	•	+	-	+
	E .	0.73	0.90	0.16	0.81	4.68	1	-	+	+
25143		3.46	0.90	1.96	3.72	1.77	1	+	-	-
7504		1.89	1.02	0.52	1.85	3.63	1	-	+	+
	71 AA405984		1.02	1.66	3.38	2.10		+	•	-
	66 AA133165	3.48	21.92	17.55	2.43	3.04	1	+	-	+
3445	1		1.12	0.86	2.37	3.09	•	-	+	-
2506	1		1.12	2.72	3.98	1.48	1	+	•	-
	10 AA449644		1.47	0.47	1.32	4.13	1	-	+	+
	29 AA489670		6.84	3.84	1.96	3.49		-	-	+
	90 AA826237		4.02		2.71	2.7		+	-	+
	118 AA010608		4.02 0.76		2.17	3.2		•	+	+
	664 H05264		0.76 2.74		2.17	3.2. 2.7		+	-	-
2519					1.14	4.3	The state of the s	•	+	+
2533					3.15	2.2		+	-	-
	232 AA121428				2.33	3.1		+	+	+
	244 AA63054				1.34	4.0	1	<u>.</u>	+	+
898.	242 AA59862	1 10.34	1.14	2.33	1.34	7.0	- 1			

395625						Table 1-3		•			
CGCAGGA ALAIT911	3056251	A	14.45	611	4 73		3.06	•	-	-	+
1031598								•	-	+	+
1031598 AA669474								•	•	+	+
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28	83034	N51280	9.61	3.05	4.60	3.15	2.09	+	+	-	-
50	64537 A	A121732	1.79	1.93	0.42	0.93	4.31	-	-	+	+
1	82734	T73651	7.21	6.90	1.72	1.05	4.18	-	-	+	+
163	31682 A	A994801	1.61	0.53	0.74	3.03	2.19	-	+		-
29	90378	N64508	17.10	17.02	4.07	1.00	4.20	-	-	+	+
20	62865	H99650	30.48	11.78	11.64	2.59	2.62	-	+	-	+ `
10:	31698 A	A609556	6.26	3.13	1.95	2.00	3.21	-	•	•	+
	09200	T81107	3.17	0.93	1.78	3.42	1.79	+	+	-	-
8:	24358 A	A489681	0.96	0.68	0.25	1.42	3.78	-	-	+	+
	32684	R20424	24.32	17.98	6.32	1.35	3.85	-	-	+	+
2	40430	H78097	3.52	1.28	1.43	2.74	2.46	+	-	-	-
		A858390	28.16	22.56	7.13	1.25	3.95	-	-	+	+
7	31272 A	A421073	5.02	2.47	1.59	2.03	3.16	-	•	-	+
	45860	N55339	30.36	12.47	11.02	2.43	2.76	-	+	+	+
8	54874 A	A630346	2.22	0.68	1.15	3.26	1.92	+	+	-	-
		A490802	1.03	0.63	0.29	1.63	3.55	-	-	-	+
14	35029 A	A857212	1.84	0.67	0.75	2.74	2.44	-	+	-	-
9	51080 A	A620446	2.43	1.63	0.66	1.49	3.69	-	-	+	+
	22731	R45264	4.46	2.73	1.26	1.64	3.54	-	-	+	+
		A447742	34.08	17.95	10.41	1.90	3.27	•	-	+	+
	258521	N40743	3.33	1.39	1.20	2.40	2.77	+	-	+	-
		AA988798	135.32	117.37	33.70	1.15	4.01	-	-	-	+
	144926	R78559	9.20	3.96	3.24	2.32	2.84	-	+	-	+
	1	AA598758	15.87	13.03	4.03	1.22	3.94	-	-	+	+
		AA478268	7.20	4.16	2.10	1.73	3.42	-	•	+	+
	1	AA877347	8.03	3.31	2.94	2.43	2.73	-	-	-	+
	277660	N49405	14.16	5.37	5.65	2.64	2.51	-	+	-	+
:	840821	AA486261	17.59	8.78	5.60	2.00	3.14	-	-	-	+
•	760344	AA426227	5.89	3.00	1.85	1.96	3.18	-	-	-	+
	813187	AA456721	22.70	10.74	7.51	2.11	3.02	-	-	-	+
	685516	AA262573	3.36	1.79	1.03	1.88	3.26	ļ -	-	-	+
	503851	AA130042	18.15	5.54	9.79	3.28	1.85	+	+	-	-
1	631762	AI025126	46.32	36.08	12.06	1.28	3.84	-	-	-	+
	343167	W67309	15.62	7.28	5.24	2.15	2.98	-	•	-	+
	30793	R18258	7.89	2.84	3.37	2.78	2.34	-	+	•	-
	82874	T69348	5.14	2.70	1.60	1.90	3.21	-	-	-	+
	209167	H63668	3.29	1.15	1.46	2.86	2.25	+	+	+	-
	811842	AA443177	9.91	4.54	3.39	2.18	2.92	-	-	-	+
	810221	AA464700	3.85	1.37	1.68	2.81	2.30	-	+	-	-
	249705	H85464	62.65	37.20	18.34	1.68	3.42	-	-	-	+
	178137	H47015	238.41	190.87	61.93	1.25	3.85	-	-	+	+
	815800	AA485216	1.22	1.44	0.29	0.84	4.25	-	-	+	+
	294259	W01800	4.21	2.21	1.32	1.90	3.19	-	-	+	+
	248454	N59626	3.70	1.95	1.16	1.89	3.19	-	-	+	-
		AA151796	50.88	17.89	22.72	2.84	2.24		•	+	•
	731257	AA416833	4.45	2.31	1.41	1.93	3.15		-	+	+
	259350	•	22.84	6.64	13.93	3.44	1.64		+	-	-
		AA894687	3.57	1.31	1.53	2.73	2.34	5	+	-	+
		AA195601	5.48	2.72	1.79	2.02	3.06		•	•	+
		AA056001	2.59	1.39	0.81	1.87	3.20	1	-	+	-
	837953	AA458578	10.44	6.85	2.94	1.52	3.55	5 -	-	+	+

					I able 1-3	_				
1500324	AA885819	6.77	4.55	1.89	1.49	3.58	•	•	+	+
167205	R90934	7.85	2.56	3.93	3.06	2.00	+	+	-	•
594693	AA165313	7.33	3.57	2.44	2.05	3.00	•	-	+	+
868119	AA634213	10.00	4.58	3.49	2.18	2.87	•	-	•	+
502463	AA134770	5.77	2.99	1.85	1.93	3.12	-	-	-	+
342271	W61215	3.90	1.09	2.69	3.60	1.45	+	+	•	-
256680	H96392	1.72	1.47	0.44	1.17	3.88	-	-	•	+
430179	AA010091	51.95	32.22	15.13	1.61	3.43	-	-	+	•
•	AA676961	10.77	5.36	3.55	2.01	3.04	-	-	•	+
25517	R11888	8.52	4.10	2.87	2.08	2.96	-	-	•	+
1632141	AI004916	2.15	0.93	0.79	2.31	2.74	-	-	-	+
	AA455714	0.98	0.77	0.26	1.28	3.77	-	- ·	+	+
	AA488588	7.02	2.32	3.49	3.03	2.01	-	+	-	-
	AA497118	0.56	0.53	0.14	1.06	3.98	-	-	•	+
42636		1.11	0.32	0.72	3.50	1.54	_	+	•	-
	AA448289	1.83	0.63	0.85	2.88	2.15	-	+	-	-
244652	N52911	24.70	12.93	7.92	1.91	3.12	-	-	+	-
	AA608583	4.17	1.22	2.62	3.43	1.59	+	+	-	-
	AA922326	13.23	7.43	4.08	1.78	3.24	•	-	+	+
	AA775431	4.20	3.49	1.10	1.20	3.82	_	-	-	+
130773		3.71	1.21	1.90	3.07	1.95	_	+	-	•
	AA129171	2.32	0.70	1.34	3.29	1.73	_	+	_	_
	AA126356	33.96	23.00	9.58	1.48	3.54	_	•	-	+
295206	B	3.16	1.64	1.02	1.92	3.10	_	_	+	+
281072	I .	· 5.02	1.92	2.10	2.62	2.39	+	-	•	
	AA482198	2.04	0.70	0.98	2.91	2.09	_	+		
143306		10.42	4.91	3.62	2.12	2.88	_	<u>.</u>	_	+
247609	1	3.15	1.20	1.33	2.63	2.37	-	+	-	, _
250678	i .	4.24	1.20	1.53	2.03	2.82			_	+
	7 AA458943	4.24 0.47	0.34	0.13	1.38	3.62	-	_	+	+
	8 AA504284	3.42	1.55	1.23	2.21	2.79	+	_	+	_
19788	•	3.42 4.97	4.63	1.27	1.07	3.92	-	_	+	+
	B C	9.31	3.39	4.15	2.75	2.25	+	_	_	_
26147 13725	1	1.35	0.58	0.51	2.35	2.64	_	+	_	+
5273	3	6.16	4.38	1.72	1.41	3.58	_	<u>,</u>	+	+
	_1			3.19	1.33	3.65	_	_		+
7625	2 T59873 3 AA598487	11.64	8.74	2.03	2.63	2.35	•	+	+	
	4 AA485214	4.77 14.69	1.81 7.77	4.77	1.89	3.08	•	7	+	+
	2 AA262559		3.70	0.90	0.97	3.99	-	-	+	+
		3.59			3.81	1.15	+	- +		•
23346		14.24	3.73	12.38		3.95	T	т.	+	+
8048		14.69	14.52	3.72	1.01		-	- +	.	
32205	3	11.58	4.27	5.15	2.71	2.25	-	т	•	+
29915		1.37	0.71	0.45	1.92	3.04	-	-	- +	+
13660	1	10.58	5.59	3.46	1.89	3.06	-	-		+
	70 AA281504	1.32	1.16	0.35	1.14	3.81	-	-	+	+
	34 AA664105	3.16	1.34	1.22	2.36	2.59	_	-	т	+
2725		6.22	4.17	1.80	1.49	3.45		-	•	+
833		5.08	3.35	1.48	1.51	3.43		-	•	+
	91 AA933888	10.79	3.85	5.05	2.80	2.14	1	+	+	+
	50 AA496628	5.88	4.08	1.68	1.44	3.50		-	₹	τ-
	55 AA459679	3.38	1.25	1.51	2.70	2.24		+	-	-
1482			1.96	1.31	1.98	2.96		-	+	-
1881	3		1.41	1.23	2.30	2.64		•	+	•
7800	09 AA478481	4.81	2.23	1.73	2.15	2.78	+	-	+	-

				•	Table 1-3					
561971	AA088886	3.81	3.31	1.01	1.15	3.78	-	-	-	+
	AA608546	3.53	2.28	1.04	1.55	3.39	•	-	+	-
266747	N31312	3.05	1.60	1.01	1.91	3.02	•	-	• .	+
129146	R10947	7.97	4.87	2.42	1.64	3.30	•	-	-	+
41940	R59087	10.51	5.11	3.65	2.05	2.88	-	•	-	+
786525	AA452113	1.46	0.42	1.00	3.46	1.46	+	-	•	•
530814	AA070226	46.55	26.81	14.60	1.74	3.19	-	-	-	+
287528	N62096	1.98	1.06	0.65	1.87	3.05	•	-	+	+
701112	AA287323	8.70	5.76	2.55	1.51	3.41	-	-	-	+
1435185	AA857632	2.56	1.69	0.75	1.51	3.40	-	-	+	+
248261	N78083	1.51	0.70	0.55	2.15	2.76	-	+	+	+
951091	AA620458	2.03	1.20	0.63	1.69	3.22	-	-	-	+
123755	R01304	5.06	2.03	2.10	2.49	2.41	+	+	+	+
505344	AA147499	2.26	1.01	0.84	2.23	2.68	-	-	-	+
827152	AA521247	3.50	2.11	1.08	1.66	3.24	•	-	+	+
812977	AA464617	7.62	5.19	2.22	1.47	3.43	-	-	+	+
1521977	AA906997	2.08	0.72	1.04	2.90	2.00	-	+	•	-
416010	W85832	53.71	32.20	16.66	1.67	3.22	-	-	+	-
1031086	AA610071	25.15	13.97	8.15	1.80	3.09	-	-	+	+
878525	AA775857	11.10	9.66	2.97	1.15	3.74	•	-	-	+
430153	AA010233	4.46	1.54	2.25	2.90	1.98	•	+	-	-
417801	W88848	17.18	13.47	4.77	1.28	3.60	-	-	+	+
108265	T70810	7.24	5.07	2.10	1.43	3.45	-	•	-	+
745296	AA625574	29.18	19.75	8.59	1.48	3.40	-	•	+	+
365098	AA025112	4.93	1.78	2.34	2.76	2.11	-	+	-	-
11278	T85958	7.75	4.81	2.38	1.61	3.25	<i>-</i>	-	-	+
81365	1 AA447761	4.96	1.93	2.17	2.58	2.29	-	+	-	-
81028	4 AA463932	2.27	1.27	0.74	1.79	3.07	-	-	+	+
72648	8 AA399264	10.10	6.03	3.17	1.68	3.18	-	-	-	+
51211	6 AA133721	4.59	1.55	2.41	2.96	1.90	+	-	+	-
81576	0 AA485018	2.31	1.17	0.81	1.98	2.87	-	-	-	+
5034	4 H17002	0.85	0.39	0.32	2.16	2.69	+	-	+	-
29383	0 N98283	4.43	1.72	1.96	2.58	2.26	+	-	+	-
21389	0 H72937	14.14	7.99	4.60	1.77	3.07	-	-	+	+
78422	25 AA446885	10.62	7.56	3.09	1.41	3.44	-	-	<u>-</u>	+
3660	7 R46837	7.52	6.30	2.06	1.19	3.65	-	-	+	+
70370	7 AA278195	4.12	2.45	1.31	1.68	3.15	-	-	•	+
400	17 R53311	8.29	4.13	2.94	2.01	2.82	-	-	-	+
6693	71 AA253444	4.99	2.55	1.74	1.96	2.87	-	-	+	-
	60 AA122048	5.94	3.29	1.97	1.81	3.02	-	•	-	+
5446	39 AA074677	2.82	1.94	0.84	1.46	3.37	-	-	-	+
4891	09 AA056484	7.44	5.33	2.17	1.40	3.43	- '	-	+	+
2793	94 N48701	3.18	1.23	1.42	2.59	2.23		-	+	-
742		5.42	3.10	1.76	1.75	3.07		-	-	+
	36 AA054643	3.54	1.49	1.46	2.38	2.43	1	-	+	_
7126	10 AA281793	1.06	0.58	0.35	1.83	2.98	1	-	-	+
4481	17 AA702480		1.64	1.61	2.38	2.43	1	+	-	+
	114 H16851		0.28	0.44	2.92	1.89	1	+	-	+
	638 AA872323		4.65	2.16	1.53	3.29		•	-	4
	399 AA487552		4.37	3.33	2.08	2.73	3	-	-	+
	799 AA664067			2.64	1.39			-	+	7
	753 AA457737			2.67	2.95		1	+	-	
	749 H23049			1.33	2.05			-	+	•
361	659 W96280	13.43	4.23	8.34	3.18	1.6	1 -	+	•	•

	252453	H87143	20.12	14.53	5.91	1.38	3.40	-	•	· +	+
	133717	R27767	9.46	2.97	5.91	3.18	1.60	+	+	-	-
		AA449037	55.68	21.05	26.01	2.65	2.14	-	+	•	-
	76196	T59668	2.53	0.90	1.29	2.83	1.96	+	+	-	-
	322561		107.98	70.00	33.31	1.54	3.24	-	-	-	+
		AA505150	1.95	1.08	0.66	1.81	2.97	-	-	-	+
	1	AA496455	4.01	2.20	1.36	1.83	2.96	-	-	-	+
	•	AA878210	8.24	4.30	2.88	1.92	2.86	-	•	+	+
	358083	W95279	2.45	0.80	1.44	3.07	1.71	-	+	-	_
		AA447731	84.51	49.31	27.59	1.71	3.06	-	•	-	+
			. 5.35	3.78	1.60	1.42	3.35	_	•	+	+
		AA455126	27.74	19.39	8.32	1.43	3.34	_	_	•	+
		AA625812	2.15	1.01	0.82	2.14	2.63	+		+	-
	321386	W44852	2.15	1.56	0.82	1.57	3.19	, -	_	_	+
	253577		1.48	1.24	0.77	1.19	3.57	_	_	_	+
	41174	R56106	4.02	2.38	1.31	1.69	3.07	_	_	_	+
	149013		4.02 7.82	2.36 5.75	2.30	1.36	3.40	_	_	+	+
	296529	1	7.82 9.83	7.09	2.92	1.39	3.37	_	_	·	+
		AA017200		3.33	2.92 1.84	1.69	3.06	<u>-</u>	_	_	+
			5.63		0.57	1.65	3.10	-	_	_	+
	154749		1.76	1.07			2.72	-	_	_	+
	197176		20.79	10.24	7.65	2.03	3.61	-	•	+	+
apple of a		AA190313	24.22	21.29	6.72	1.14		•	+	•	•
AND THE PARTY OF T		AA598561	14.33	5.93	6.17	2.42	2.32	-	+	-	+
4.		AA147837	10.43	6.02	3.47	1.73	3.00	-	-	-	+
		AA042928	9.14	5.47	2.99	1.67	3.06	-	•	-	т
		AA872122	85.21	65.01	24.92	1.31	3.42	-	-	+	-
	283196	1	3.80	2.01	1.34	1.89	2.84	-	-	+	-
		AA126708	18.65	13.11	5.65	1.42	3.30	-	-	+	+
		AA873060	6.33	4.83	1.85	1.31	3.41	-	-	+	+
		1 AA479154	6.80	4.46	2.13	1.53	3.20	-	-	+	+
55	325600	1	7.30	3.27	2.93	2.23	2.49	-	•	-	+
To a supplement of the supplem		9 AA455917	14.02	6.63	5.40	2.11	2.60	-	•	-	+
Fore	3207	5	1.03	0.37	0.54	2.79	1.92	-	+	-	-
TOTAL STATE OF THE PARTY OF THE		0 AA777781	7.53	4.62	2.45	1.63	3.08	-	-	-	+
Contract Con		0 AA962236	6.44	4.29	2.01	1.50	3.21	-	-	+	+
2		7 AA664155	8.56	4.10	3.27	2.09	2.62	-	-	-	+
2 2		2 AA677185	7.68	5.14	2.40	1.49	3.20	-	-	+	+
	2661		1.94	0.59	1.40	3.31	1.39	-	+	-	-
		9 AA988701	24.27	24.53	6.55	0.99	3.70	-	•	-	+
	34272		14.93	6.70	6.06	2.23	2.46	-	-	+	+
		IS AA156202	4.37	5.11	1.14	0.85	3.84	-	-	+	+
		50 AA488481	5.13	1.90	2.57	2.69	1.99	-	+	-	-
	28228	1	0.84	1.00	0.22	0.84	3.85	-	-	-	+
	20313	R .	12.63	11.94	3.48	1.06	3.63	•	•	+	+
	4033		2.61	1.86	0.79	1.40	3.28		-	-	+
	2041	l l	2.40	1.26	0.86	1.90	2.78		-	+	-
		83 AA431407	1.32	0.85		1.55	3.13		-	+	-
	1419			10.40		1.82	2.86		-	-	+
	3568	8		0.94		3.38	1.30		+	-	-
		66 AA033564		1.68		1.62	3.05	E	-	-	+
	2466			15.47		1.09			•	•	+
		21 AA485688		1.34		0.80			-	•	+
		32 AA284337		2.02		1.61			-	-	+
	426	59 R61126	10.21	3.36	6.30	3.04	1.62	-	+	-	•

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4	460841	AA708189	1.02	0.58	0.35	1.75	2.91	•	•	-	+
	306771	N91887	3.61	1.62	1.49	2.23	2.42	•	-	-	+
	160485	H22136	5.17	2.96	1.78	1.74	2.90	-	-	+.	+
•	950461	AA599107	38.86	34.73	11.02	1.12	3.53	•	-	+	+
	75886	T59478	14.08	9.92	4.36	1.42	3.23	•	-	+	+
	283919	N50797	7.56	4.94	2.43	1.53	3.11	-	-	+	+
	813271	AA455938	0.96	1.12	0.25	0.85	3.79	-	-	-	+
	823627	AA490561	3.25	1.81	1.14	1.79	2.85	-	-	-	+
	242706	H93552	5.15	2.96	1.78	1.74	2.90	-	-	-	+
	970271	AA775957	3.16	1.23	1.52	2.56	2.07	-	+	-	-
	783645	AA446690	6.96	2.63	3.51	2.65	1.98	•	+	-	-
	486459	AA042858	6.24	5.89	1.75	1.06	3.57	-	-	+	+
	884346	AA629524	5.38	3.61	1.71	1.49	3.14	-	-	-	+
	26406	R20638	6.56	5.37	1.92	1.22	3.41	-	-	-	+
1	467108	AA883160	4.29	2.67	1.42	1.61	3.02	•	-	-	+
1	632161	B .	39.75	38.39	11.08	1.04	3.59	-	-	-	+
		AA599120	9.36	3.14	5.71	2.98	1.64	-	+	-	-
		AA463200	9.64	6.52	3.08	1.48	3.13	•	•	-	+
		AA669183	4.20	2.32	1.50	1.81	2.80	-	-	+	+
		AA456271	11.89	6.59	4.25	1.80	2.80	•	•	•	+
		AA074224`	2.79	1.08	1.39	2.58	2.02	-	+	-	-
		AA699770	3.42	4.00	0.91	0.85	3.74	-	-	+	+
	324342		4.41	4.29	1.24	1.03	3.57	•	-	-	+
		AA029295	3.09	1.27	1.44	2.44	2.15	-	+	-	+
		AA489343	28.44	25.71	8.17	1.11	3.48	•	-	-	+
	129725	1	17.94	10.53	6.22	1.70	2.88	•	-		+
	12119		22.80	22.77	6.36	1.00	3.58	-	•	+	+
		AA457235	3.58	4.32	0.95	0.83	3.76	-	-	+	++
	28314		3.58	1.94	1.31	1.84	2.74	-	-	-	+
	163670	1	19.22	12.06	6.43	1.59	2.99	-	-	•	Ţ
	12398	L .	2.38	0.92	1.19	2.59	1.99	-	+	-	-
	25300	1	115.86	35.75	86.54	3.24	1.34	+	+	-	- +
		9 AA705058	0.92	0.61	0.30	1.51	3.07	-	-	+	
		5 AA630584	1.30	0.82	0.43	1.57	3.00	-	-	+	+
	5115		3.26	3.41	0.90	0.96	3.61 3.57	•	-	<u>.</u>	+
		3 AI732248	1.22	1.22	0.34	1.00	2.71	<u>-</u>	_	_	+
		0 AA455931	4.14	2.24	1.53	1.85	2.71	<u>-</u>	<u>-</u>	_	+
	21120	3	4.71	2.93	1.60 10.07	1.61 1.15	3.41	-	-	-	+
		1 AA412053	34.32	29.87		2.47	2.08		+	_	+
		0 AA873577	35.19	14.23	16.93	2.47	2.52	+		+	· -
		38 AA197344	7.37	3.62	2.93 1.79	1.52	3.03	_	_		+
	2892	1	5.44	3.58 1.79	1.79	1.83	2.72		_	+	
		25 AA457717		5.76	3.20	1.62	2.72		•	+	
	2430	1	9.34	5.82	3.51	1.71	2.83		_	+	
		26 AA430615		5.73	2.31	1.71	3.23		_	•	+
	2898 1963			41.22	79.12	2.98	1.55		+	-	-
		45 K92455 80 AA465424		41.22	3.29	2.98	2.49		•	+	•
	745	t t		3.37	1.75	1.55	2.98	_	•	•	+
		13 AA412643		2.39	1.73	1.65	2.88	-	•	-	+
	7303 348			2.06	0.85	1.33	3.20	1	•	+	-
		74 AA608514		11.69	14.05	2.47	2.06		+	-	-
	778	4		9.75		1.59	2.93	7	-	-	+
		04 AA894577		2.66		1.59	2.93	2	-	-	+
	4 -1 / 20-	- 10 4 307 707 7			,	3.20		•			

	263070	N20052	1.17	0.90	0.37	1.31	3.21	-	•	•	+
	951216	AA620565	2.02	1.34	0.67	1.50	3.02	-	-	-	+
	308726	N93236	14.73	5.64	7.73	2.61	1.91	-	+	•	•
	40136	R53348	5.23	1.99	2.77	2.62	1.89	+	-	•	-
	1461528	AA883711	3.95	2.84	1.27	1.39	3.12	•	-	+	+
	760148	AA424344	2.93	1.10	1.58	2.66	1.85	•	+	•	•
	430235	AA010223	3.43	1.72	1.36	1.99	2.52	•	-	•	+
	884867	AA669443	24.71	16.78	8.14	1.47	3.03	-	-	•	+
	744944	AA625890	1.73 ·	0.90	0.67	1.92	2.59	-	-	•	+
	202904	H54020	12.77	10.17	3.93	1.26	3.25	-	-	-	+
	257382	N30699	12.80	10.78	3.86	1.19	3.32	-	-	+	+
	590390	AA157150	6.42	3.41	2.45	1.88	2.62	•	-	•	+
	460798	AA708152	1.30	0.65	0.52	2.00	2.50	-	-	-	+
	815164	AA481135	0.88	0.71	0.27	1.25	3.25	-	-	-	+
	214884	H74119	3.55	2.09	1.27	1.70	2.79	-	-	-	+
	782439	AA431836	35.89	20.99	12.92	1.71	2.78	-	-	-	+
	277596	N57002	2.91	1.45	1.17	2.00	2.49	-	-	+	-
	344432		15.48	11.50	4.93	1.35	3.14	-	-	-	+
	796341	AA459750	7.16	4.60	2.44	1.56	2.93	-	-	-	+
	731308	AA416759	4.84	1.70	2.96	2.85	1.64	•	+	-	-
	788518	AA452566	13.97	9.65	4.61	1.45	3.03	-	-	-	+
	201628	R98008	7.51	5.63	2.39	1.33	3.14	•	-	+	+
THE PROPERTY OF THE PROPERTY O	1336262	AA872057	35.80	21.64	12.69	1.65	2.82	-	-	-	+
		AA598595	1.79	2.04	0.50	0.88	3.59	•	-	-	+
and the second		AA497010	15.69	7.46	6.62	2.10	2.37	-	+	-	+
A CALL		AA099855	33.49	26.22	10.49	1.28	3.19	-	-	+	+
		AA453713	0.52	0.34	0.18	1.52	2.95	-	-	-	+
222		AA430205	2.67	1.22	1.18	2.19	2.27	•	-	+	-
E-102	1492967	AA886333	3.68	1.85	1.49	1.99	2.47	+	-	+	-
	31225	1	4.76	1.64	3.09	2.90	1.54	-	+	-	-
	35807	I .	2.91	1.90	1.00	1.53	2.91	-	-	-	+
print grant printerprinter printerprinter printerprinter printerprinter printerprinter	950450	AA599094	1.77	1.02	0.66	1.75	2.70	-	-	-	+
Count E s	81494	5 AA465116	4.97	2.92	1.82	1.70	2.74	-	-	-	+
2.5		5 AA453926	13.21	6.75	5.32	1.96	2.48	-	-	-	+
parties a partie of the parties of t	78909	1 AA452933	9.50	6.97	3.09	1.36	3.08	-	-	+	+
10000	84314	0 AA486516	14.94	10.35	4.99	1.44	3.00	-	-	-	+
		1 AA456975	52.82	30.51	19.51	1.73	2.71	-	•	•	+
i i	87840	3 AA670353	7.20	6.82	2.13	1.06	3.38	-	-	+	+
	86883	8 AA775223	3.29	2.14	1.14	1.54	2.90	-	-	+	-
	22252	7 H84287	2.88	2.10	0.94	1.37	3.05	-	-	-	+
	84276	7 AA486300	2.15	1.48	0.73	1.46	2.96	-	-	-	+
	89767	0 AA496800	6.35	2.19	4.18	2.90	1.52	-	+	-	•
	74258	31 AA400247	1.20	1.08	0.36	1.11	3.31	-	-	-	+
	5233	8 H23277	2.91	1.24	1.41	2.35	2.07	-	+	-	+
	59039	98 AA148034	12.27	6.94	4.63	1.77	2.65	-	-	•	+
	84062	20 AA488036	13.81	8.57	4.92	1.61	2.81	-	-	-	+
	85489	99 AA630374	18.74	13.26	6.24	1.41	3.00	-	-	-	+
	382	53 R35943	2.80	1.95	0.94	1.44	2.98	-	-	-	+
		69 AA156940	1.41	0.77	0.55	1.82	2.59	-	-	-	+
	3436			0.58	0.94	2.73	1.68	-	+	-	-
		96 AA286914		1.54	0.78	1.48	2.93	-	•	+	+
		52 AA886870		11.50	2.79	0.86	3.54	-	-	+	+
	2033	8		0.74	1.05	2.57	1.83	-	+	-	-
	4517	07 AA707659		5.80	2.45	1.31	3.09	-	-	+	+

1			21.66		1 10	2 22 1			_	+
22747	T75072	37.20	31.66	11.55	1.18	3.22	•	•	<u>.</u>	,
	AA465180	3.49	1.97	1.33	1.77	2.62	-	•	. +	+
251731	H96904	0.82	0.84	0.24	0.98	3.41	•	•	-	
	AA456833	4.86	3.88	1.55	1.25	3.13	•	-	•	+
	AA190747	1.58	1.06	0.55	1.49	2.88	-	-	•	+
39821	R53428	5.84	5.87	1.73	0.99	3.38	•	-	•	+
898210	AA598597	8.13	8.02	2.42	1.01	3.35	-	-	-	+
46367	H09959	1.63	0.62	0.94	2.62	1.74	+	•	-	•
28436	R40663	0.89	0.37	0.46	2.43	1.93	+	-	-	-
259953	N32594	1.46	1.97	0.40	0.74	3.63	-	-	-	+
307882	W21373	7.09	4.18	2.66	1.69	2.67	-	-	-	+
	AA283029	0.89	0.56	0.32	1.58	2.78	-	-	-	+
195903	R92227	6.95	5.56	2.24	1.25	3.11	-	-	-	+
42076	R60847	5.72	3.44	2.12	1.66	2.69	•	-	•	+
	AA887201	2.03	1.50	0.68	1.36	3.00	-	-	-	+
	AA128297	51.40	43.89	16.15	1.17	3.18	-	-	-	+
	AA457517	2.51	1.48	0.94	1.69	2.66	-	-	_	+
281191		7.36	3.32	3.45	2.22	2.13	_	+	•	+
	N50983	4.56	3.32 2.74	1.70	1.66	2.68	_	· -	_	+
391984			4.14	1.65	1.24	3.11	_	-	+	+
1635665		5.13	0.37	0.19	1.46	2.89	_	_	•	+
49485		0.54	3.27	1.98	1.63	2.71	_	_	_	+
	AA033973	5.35				2.93	_	_	_	+
	AA043347	29.51	20.98	10.06	1.41		-	_	_	+
	AA287948	1.54	1.24	0.50	1.23	3.10	•	-	-	+
144977	1	71.81	48.67	25.09	1.48	2.86	•	-	-	_
240961	1	2.11	0.82	1.20	2.57	1.76	-	+	•	+
	AA778675	3.07	1.85	1.15	1.66	2.67	-	-	-	
45515	3	3.88	2.73	1.33	1.42	2.91	-	-	+	+
	AA446743	0.26	0.10	0.15	2.61	1.72	-	+	-	-
	AA443152	0.44	0.47	0.13	0.93	3.40	-	•	-	+
41597		24.58	18.77	8.16	1.31	3.01	-	-	-	+
6863	6 T49801	4.83	2.97	1.80	1.63	2.69	-	-	-	+
3581	2 R14631	2.29	2.07	0.71	1.11	3.21	-	-	-	+
56484	6 AA 129297	1.90	1.12	0.72	1.69	2.63	-	-	•	+
50309	7 AA151486	7.27	4.27	2.78	1.70	2.61	-	-	+	-
13911	3 R63022	4.32	3.44	1.41	1.26	3.06	-	-	+	+
13162	1 R23790	10.21	3.63	6.81	2.82	1.50	-	+	•	-
15638	•	2.14	1.88	0.67	1.14	3.18	-	-	-	+
78426	4 AA447021	5.67	3.30	2.18	1.72	2.60	-	-	-	+
29170		2.41	1.00	1.27	2.41	1.90	+	-	+	-
31049		7.56	7.25	2.32	1.04	3.26	-	-	-	+
	6 AA443698	33.80	23.65	11.75	1.43	2.88	-	-	-	+
28214	1	30.16	18.11	11.43	1.67	2.64	-	-	+	-
	11 AA700736	1.73	1.11	0.63	1.56	2.74	-	-	•	+
25290	1		2.26	2.83	2.39	1.91		+	-	•
	73 AA701655		9.66	3.34	1.11	3.19	-	-	-	+
	41 AA406040		3.98	2.43	1.63	2.67		-	-	+
	71 AA598868		7.34	4.24	1.57	2.73		-	-	+
	68 AA521292		2.85	3.24	2.29	2.01	L .	-	+	-
1413			0.68		1.90	2.40	1	-	+	-
	98 AA418383		16.76		1.16	3.14	•	•	-	+
	06 AA701981		2.42		1.71	2.58	1	-	-	+
	88 AA629554		3.97		1.23	3.06		-	-	+
462	1		0.35		2.97	1.31	E .	+	-	-
702	7 11 1107132	. 1.03	0.00	29			1			

							1				
	253253	H89104	1.73	1.30	0.58	1.33	2.95	•	-	-	+
		AA609106	3.21	2.11	1.16	1.52	2.76	-	-	•	+
	1	AA011445	3.77	1.23	3.11	3.06	1.21	-	+	•	-
	52327	H23459	5.70	4.77	1.85	1.19	3.08	•	-	•	+
		AA457725	72.43	48.42	26.10	1.50	2.78	-	•		+
	155838	R72215	2.53	1.96	0.85	1.29	2.97	-	-	+	•
	135083	R33917	17.07	17.59	5.18	0.97	3.29	•	-	•	+
	271472	N35020	11.80	7.84	4.28	1.50	2.76	-	-	-	+
	29964	R14699	1.12	0.63	0.45	1.79	2.47	-	-	-	+
	269300	N35863	10.64	7.57	3.73	1.41	2.85	-	•	-	+
	429678	AA011593	3.12	0.86	4.85	3.61	0.64	+	+	-	-
	898096	AA598795	4.06	2.82	1.45	1.44	2.81	•	-	-	+
	730554	AA436031	11.89	8.38	4.20	1.42	2.83	-	-	•	+
	260628	H97565	8.94	7.49	2.93	1.19	3.05	-	-	-	+
	730412	AA469966	1.25	1.00	0.42	1.25	3.00	-	-	-	+
	825771	AA491312	12.13	9.01	4.19	1.35	2.90	-	-	+	-
	342211	W63789	1.92	1.55	0.64	1.24	3.00	•	-	•	+
	845609	AA644345	15.29	12.54	5.07	1.22	3.02	-	•	-	+
	814154	AA496252	2.29	1.38	0.89	1.67	2.57	-	-	+	-
	897730	AA598987	41.12	27.32	15.07	1.50	2.73	-	-	•	+
	141336	R63823	2.12	1.03	0.97	2.05	2.18	+	-	+	-
	263894	N28522	0.46	0.42	0.15	1.10	3.13	-	-	-	+
	252864	H88424	1.20	1.00	0.40	1.20	3.03	-	-	-	+
	291057	1	1.34	0.61	0.66	2.18	2.04	-	+	-	-
17,111		AA417805	6.05	3.90	2.27	1.55	2.67	-	-	-	+
200	132911	1	52.56	22.96	27.26	2.29	1.93	-	+	-	+
and the		AA099404	2.11	1.57	0.73	1.34	2.87	-	-	-	+
		AA678065	2.38	1.00	1.30	2.39	1.83	-	+	-	-
1111		AA947922	4.13	3.78	1.32	1.09	3.12	-	-	-	+
		7 AA464995	0.44	0.43	0.14	1.04	3.16	-	-	-	+
20		9 AA495962	4.17	3.13	1.45	1.33	2.87	-	-	-	+
and designed	27107	1	4.31	3.13	1.53	1.38	2.82	-	-	-	+
		8 AA424045	4.05	3.18	1.39	1.27	2.91	-	-	+	-
FRI		1 AA070997	5.75	3.77	2.16	1.52	2.66	-	-	-	+
		8 AA457082	2.82	2.95	0.87	0.96	3.23	-	-	-	+
gazet.		3 AA161283	0.66	0.64	0.21	1.04	3.14	-	-	-	+
117011	3638		3.19	3.07	1.02	1.04	3.14	-	-	-	+
		4 AA599099	9.75	5.67	3.98	1.72	2.45	+	-	+	-
		7 AA598625	2.02	1.65	0.68	1.22	2.95	-	-	-	+
-		7 AA463449	1.59	1.01	0.61	1.57	2.59	-	-	-	+
	27204	3	6.87	3.74	2.95	1.84	2.33	-	-	-	+
	32861		0.46	0.40	0.15	1.13	3.03	-	-	-	+
		88 AA705047	0.55	0.43	0.19	1.27	2.89	-	_	-	+
		02 AA989139	2.25	1.64	0.81	1.37	2.79	-	-	-	+
		02 AA287936	4.23	3.51	1.43	1.20	2.96	-	_	+	+
	2343		3.16	2.04	1.21	1.55	2.61	_	-	-	+
	2516		0.50	0.47	0.16	1.06	3.09	_	-	-	+
	508		7.95	7.40	2.58	1.08	3.08		-	-	+
		35 AA521108	2.24	1.62	0.81	1.39	2.77		-	•	+
	3029		10.63	10.42		1.02		1	-	+	+
		38 AA872704	238.01	171.23		1.39		1	•	-	+
		94 AA669452		28.65		1.45			-	-	+
		79 AI016074		2.02		1.62		1	-	•	+
	2021			16.52		1.78		E .	-	+	-
	2021	2.1 1152505	22.40	2	.20	1.,0	2.50	1			

80764 T63027

825012 AA489194

3.05

26.33

8.71

1.76

26.16

4.46

					Table 1-3					
197821	R93744	5.95	4.32	2.15	1.38	2.77	-	-	+	+
951022 A		3.02	1.90	1.18	1.59	2.55	-	•	-	+
856535 A		1.40	0.50	1.04	2.79	1.35	-	+	• ,	-
	A487020	2.47	3.20	0.73	0.77	3.36	-	-	•	+
197765	R93767	8.83	3.78	4.92	2.34	1.79	+	•	-	-
125308	R05886	2.61	0.93	1.97	2.80	1.33	-	+	•	-
323611	W44508	1.25	0.50	0.76	2.48	1.64	•	+	-	•
	A668897	12.39	4.81	8.02	2.58	1.55	-	+	-	-
3	AA629591	9.95	9.24	3.27	1.08	3.04	-	-	+	+
813712	AA453849	7.62	7.57	2.45	1.01	3.11	•	•	+	+
811097	AA485675	2.59	1.28	1.23	2.02	2.10	+	•	+	•
243321	H95088	2.79	1.90	1.05	1.47	2.65	•	•	+	•
856575	AA633658	2.75	2.77	0.88	0.99	3.12	•	•	+	+
360232	AA013099	2.20	1.94	0.74	1.13	2.98	-	•	-	+
26185	R20616	1.15	0.80	0.43	1.43	2.69	-	-	-	+
469685	AA027912	4.95	4.33	1.67	1.14	2.97	-	•	-	+
841501	AA487382	5.94	4.80	2.07	1.24	2.88	-	-	•	+
428215	AA001745	42.11	30.60	15.40	1.38	2.73	-	•	•	+
825356	AA504492	1.18	0.90	0.42	1.32	2.79	•	•	-	+
884480	AA629719	11.68	10.42	3.91	1.12	2.99	-	-	-	+
813179	AA456717	6.60	2.40	4.88	2.75	1.35	+	-	•	-
645368	AA206267	8.48	7.77	2.82	1.09	3.01	-	-	•	+
666371	AA232645	0.85	1.07	0.26	0.79	3.30	-	•	-	+
768179	AA426629	3.80	3.11	1.32	1.22	2.87	-	-	•	+
1420830	AA826324	3.06	2.56	1.06	1.19	2.90	-	-	-	+
246541	N77514	10.77	8.71	3.78	1.24	2.85	-	-	-	+
742695	AA401521	1.83	1.55	0.63	1.18	2.90	-	-	-	+
451769	AA706795	5.25	5.71	1.66	0.92	3.16	-	-	+	+
827132	AA521232	6.63	4.13	2.68	1.60	2.48	-	-	-	+
460218	AA677457	2.10	1.12	0.95	1.87	2.21	-	-	-	+
40075	R52015	3.13	3.07	1.03	1.02	3.05	-	-	-	+
127928	R08932	3.67	2.74	1.35	1.34	2.73	-	-	-	+
279785	N48356	0.91	0.66	0.34	1.38	2.68	-	-	+	-
1591622	AA983467	49.35	42.75	16.96	1.15	2.91	-	-	-	+
824640	AA482150	6.96	6.37	2.35	1.09	2.97	•	•	-	+
1635350	AI014387	10.17	8.91	3.48	1.14	2.92	-	-	+	-
855723	AA663960	6.73	7.60	2.12	0.89	3.17	-	-	•	+
1475746	AA873762	12.48	10.32	4.39	1.21	2.84	-	•	•	+
845502	2 A1791165	3.87	3.96	1.26	0.98	3.07	-	•	-	+
897983	3 AA598874	26.99	22.87	9.42	1.18	2.86	-	•	-	+
	6 AA490734	4.08	3.05	1.51	1.34	2.70	-	•	+	-
62768	7 AA196287	56.19	60.68	18.11	0.93	3.10	-	•	-	+
3288	7 R43699	3.04	2.20	1.15	1.39	2.64	-	-	<u>-</u>	+
	0 AA700604	28.82	38.07	8.83	0.76	3.26	-	-	+	-
81298	9 AA464623	2.18	1.75	0.79	1.25	2.77	-	-	•	+
24415		4.94	3.91	1.79	1.26	2.76	-	•	•	+
27868		13.10	9.87	4.88	1.33	2.69	-	-	-	+
32253		3.18	2.15	1.26	1.48	2.53	-	-	+	-
	2 AA733038	6.74	4.35	2.74	1.55	2.46	-	•	•	+
	70 AA669124	4.11	2.43	1.78	1.69	2.31	-	•	-	+
	26 AA856556		28.07	10.55	1.09	2.91	-		-	+
14758	31 AA872153	14.80	6.17	9.24	2.40	1.60		+	-	- -

3.00

2.53

0.99

1.46

Page 19

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51981 H23422 19.62 13.04 7.89 1.50 2.49	+
1591264 AA955007 3.18 2.86 1.11 1.11 2.88	+
898265 AA598965 20.16 18.09 7.03 1.11 2.87	+
511459 AA115310 37.04 46.42 11.64 0.80 3.18	+
296429 W00890 5.99 4.85 2.18 1.23 2.75	+
754449 AA410480 1.78 1.58 0.63 1.13 2.85	+
813149 AA456298 6.11 6.08 2.05 1.00 2.97 +	+
741988 AA402915 1.74 1.20 0.69 1.45 2.52	+
825842 AA504772 6.00 6.39 1.98 0.94 3.03 +	+
1492412 AA878561 236.26 205.84 83.73 1.15 2.82	+
1422794 AA827405 2.62 1.78 1.05 1.47 2.49	+
731104 AA421455 0.65 0.64 0.22 1.02 2.93	+
80292 T64433 2.18 1.64 0.83 1.33 2.62	+
969844 AA663826 9.12 11.76 2.88 0.78 3.17 +	+
75059 T50370 21.82 10.94 11.17 2.00 1.95 - + -	+
303139 W19519 2.17 1.22 1.00 1.77 2.17 +	-
48033 H11760 0.51 0.50 0.17 1.02 2.92	+
282720 N50079 8.31 6.24 3.18 1.33 2.61	+
291974 N73091 57.24 40.53 22.63 1.41 2.53	+
307029 N89671 56.38 44.33 21.22 1.27 2.66	+
593520 AA165402 6.25 6.33 2.13 0.99 2.93	+
1471829 AA873351 99.54 79.29 37.33 1.26 2.67 +	-
461955 AA779968 1.94 0.91 1.09 2.13 1.78 - + -	-
27072 R37780 0.29 0.27 0.10 1.07 2.85	+
282895 N51226 3.91 1.96 2.03 1.99 1.93 + - +	-
866866 AA679341 3.00 2.04 1.23 1.47 2.45	+
878373 AA670330 31.06 35.83 10.19 0.87 3.05	+
471641 AA034945 4.79 2.74 2.22 1.75 2.16 - + -	+
447568 AA702422 9.00 7.57 3.31 1.19 2.72	+
866276 AA663160 1.94 1.83 0.68 1.06 2.84	+
246800 N59078 6.67 7.15 2.25 0.93 2.96	+
565062 AA133896 1.38 1.65 0.45 0.84 3.05	+
42313 R60946 1.33 1.07 0.51 1.24 2.64	+
753917 AA479100 6.49 5.74 2.36 1.13 2.75	+
234036 H68988 4.77 3.83 1.81 1.24 2.64	+
884333 AA629528 4.10 2.83 1.69 1.45 2.43	+
758347 AA403072 2.75 2.38 1.01 1.16 2.72	+
795738 AA460286 2.06 1.70 0.77 1.21 2.66	+
72616 T51995 1.67 1.29 0.65 1.29 2.58	+
251753 H96908 2.29 3.06 0.74 0.75 3.12 +	-
950607 AA608548 19.79 11.09 9.50 1.78 2.08 +	-
825859 AA504784 1.64 1.22 0.65 1.35 2.52	+
161456 H25546 13.35 10.78 5.08 1.24 2.63 +	-
897577 AA496881 0.52 0.50 0.18 1.03 2.82	+
1599489 AA961361 5.51 4.50 2.10 1.22 2.62	+
195652 R89403 18.03 15.86 6.65 1.14 2.71	+
431597 AA676422 5.45 2.95 2.73 1.85 2.00 + - +	-
486340 AA044123 16.47 11.82 6.73 1.39 2.45	+
742806 AA400362 1.76 2.10 0.59 0.84 2.99	+
770935 AA433877 86.21 50.78 40.44 1.70 2.13	+
53165 R16153 0.89 0.87 0.32 1.02 2.80	+
307231 N93428 2.04 1.57 0.81 1.30 2.53 +	•
826459 AA521083 12.36 7.59 5.65 1.63 2.19	+
868308 AA634008 65.99 60.02 24.35 1.10 2.71	+

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	1056203	AA621031	1.92	0.89	1.16	2.15	1.65	+	-	•	-
	486175	AA043133	10.39	9.02	3.91	1.15	2.66	-	-	+	-
	81484	T63342	10.87	5.15	6.40	2.11	1.70	•	+		•
	429047	AA007509	3.17	1.48	1.90	2.14	1.67	•	+	•	-
	783697	AA446839	3.19	2.89	1.18	1.10	2.70	-	-	•	+
	784128	AA446759	1.83	1.82	0.65	1.00	2.80	•	-	+	-
	253865	N22007	1.68	1.49	0.63	1.13	2.67	-	•	+	+
	1291658	AA776810	2.05	1.00	1.17	2.05	1.75	-	+	-	-
	595604	AA167269	9.41	8.19	3.56	1.15	2.65	•	-	-	+
	784129	AA432085	6.53	5.53	2.50	1.18	2.61	-	-	+	-
	251452	H97993	3.12	2.79	1.17	1.12	2.67	•	-	+	-
	767346	AA418557	4.50	3.43	1.82	1.31	2.47	-	-	+	-
	138865	R62742	5.42	4.35	2.14	1.25	2.53	-	-	-	+
	770983	AA427401	7.24	6.59	2.70	1.10	2.68	•	-	+	-
	43705	H05706	0.35	0.33	0.13	1.07	2.71	-	-	-	+
	76005	T47291	2.35	1.80	0.95	1.31	2.46	-	-	-	+
	156045	R72518	2.01	1.50	0.83	1.34	2.43	-	-	-	+
	784085	AA446639	3.93	3.17	1.56	1.24	2.52	-	-	-	+
	418053	W90732	7.57	7.14	2.80	1.06	2.70	•	•	•	+
	322192	W37779	7.97	9.95	2.69	0.80	2.96	-	-	+	+
	309864	1	2.49	1.62	1.12	1.54	2.22	+	-	+	•
	44975	1	8.91	10.72	3.05	0.83	2.92	-	-	-	+
	825719	AA504834	0.76	0.67	0.29	1.14	2.62	-	-	-	+
	291394		6.46	4.22	2.92	1.53	2.22	•	-	+	-
:		AA461311	0.78	0.80	0.28	0.98	2.77	-	-	•	+
		AA480820	2.46	1.75	1.05	1.40	2.35	-	-	-	+
		AA045792	3.14	2.24	1.34	1.40	2.34	-	-	-	+
		AA629808	156.92	121.14	64.25	1.30	2.44	-	-	-	+
	41411	1	112.58	99.23	43.25	1.13	2.60	-	-	•	+
	247546	L	11.16	8.98	4.48	1.24	2.49	-	-	+	-
		1 AA496173	3.75	3.51	1.41	1.07	2.67	-	-	•	+
	3766	1	1.01	0.72	0.43	1.40	2.33	-	-	-	+
	56345	1 AA113347	4.17	1.82	2.90	2.29	1.44	-	+	-	-
	72550	3 AA292995	4.55	3.15	1.99	1.45	2.28	-	-	-	+
	152279	9 AA902249	1.23	1.56	0.42	0.79	2.94	-	-		+
	26817	1	28.02	27.47	10.37	1.02	2.70	-	-	-	+
		2 AA496148	5.14	3.76	2.18	1.37	2.36	-	-	+	-
		8 AA857413	249.73	233.03	94.34	1.07	2.65	-	-	-	+
		0 AA487934	4.05	3.61	1.56	1.12	2.60	-	-	-	+
	95098	9 AA620404	2.48	1.95	1.02	1.28	2.44	-	-	-	+
		6 AA481438	17.53	6.47	17.49	2.71	1.00	+	-	-	-
		6 AA634164	6.77	5.15	2.83	1.32	2.39	-	-	+	+
		31 AA633768	54.39	52.93	20.36	1.03	2.67	-	-	-	+
	155552	23 AA974971	1.21	0.85	0.53	1.41	2.28	-	-	+	-
	20678	1	6.05	5.71	2.30	1.06	2.63	-	-	-	+
	7810	61 AA446479	0.73	0.44	0.35	1.64	2.05	-	-	-	+
	3450:		0.87	0.35	0.71	2.47	1.22	. -	+	-	•
		25 AA496736	2.58	3.15	0.90	0.82	2.87	' -	-	•	+
		64 AA046321	2.58	2.44	0.98	1.06	2.63	-	-	-	+
		36 AA778869		4.43	2.04	1.16	2.52	-	-	-	+
	766	L		2.10	0.99	1.18	2.50) -	-	•	+
		32 AA668531	2.57	2.48	0.97	1.04	2.64	1	•	-	+
		95 AA454862		2.07	1.21	1.35	2.32		-	•	+
	6321	37 AA167728	0.80	0.57	0.35	1.41	2.27	7 -	-	-	+
		•									

122183	T98628	7.61	10.97 .	2.56	0.69	2.98	-	•	•	+
284545	N64762	1.10	0.77	0.49	1.42	2.24	-	•	-	+
	AA485024	1.44	1.12	0.61	1.29	2.38	-	•	-	+
629498	AA192765	3.73	2.69	1.64	1.39	2.28	•	-	-	+
243784	N33927	2.99	2.12	1.33	1.41	2.25	-	-	-	+
	AA457529	41.97	41.10	15.95	1.02	2.63	-	-	-	+
	AA704459	1.29	1.51	0.46	0.86	2.79	-	•	-	+
83011	T70522	32.97	35.26	12.18	0.93	2.71	-	-	-	+
	AA679352	10.73	11.79	3.94	0.91	2.73	•	-	-	+
277761	N49605	2.43	1.65	1.12	1.47	2.17	•	-	•	+
	AA099593	4.03	4.49	1.47	0.90	2.74	-	-	-	+
	AA463822	1.35	1.17	0.55	1.15	2.48	-	-	+	-
33045		0.63	0.43	0.29	1.47	2.15	-	-	+	-
137984	E .	1.63	1.42	0.66	1.15	2.47	-	-	. +	-
	AA968896	3.08	2.54	1.28	1.21	2.40	-	• *	-	+
138861	1	1.72	1.35	0.74	1.27	2.33	-	•	•	+
289645	l .	0.92	0.67	0.42	1.38	2.21	•	-	+	-
140792		1.66	0.85	1.02	1.96	1.63	+	-	-	-
130136	1	0.29	0.22	0.12	1.29	2.29	-	-	+	-
	AA131466	3.64	4.28	1.33	0.85	2.74	-	-	+	+
	AA521243	2.52	2.65	0.96	0.95	2.63	-	-	•	+
	AA487650	11.96	14.60	4.33	0.82	2.76	-	-	-	+
	AA599178	12.52	17.40	4.38	0.72	2.86	-	-	-	+
450464	AA682819	3.97	2.31	2.13	1.72	1.86	-	-	-	+
743211	AA400214	3.84	1.71	2.87	2.24	1.34	+	-	-	-
460164	AA676920	0.80	0.68	0.33	1.17	2.41	-	-	-	+
856174	AA630620	0.25	0.18	0.11	1.36	2.21	•	-	-	+
142320		1.49	0.78	0.90	1.92	1.65	+	-	-	-
	2 AA872143	224.84	89.33	214.43	2.52	1.05	+	-	-	-
	5 AA420992	2.04	1.38	0.98	1.48	2.09	-	-	+	-
	9 AA677499	14.78	16.78	5.51	0.88	2.68	-	-	-	+
24420	1	9.25	9.97	3.51	0.93	2.63	-	-	-	+
	8 AA456931	53.19	53.66	20.71	0.99	2.57	-	-	-	+
	8 AA774230	6.47	4.45	3.08	1.45	2.10	-	-	-	+
25141		0.69	0.68	0.27	1.01	2.52	-	~	•	+
	1 AA598675	2.02	3.02	0.71	0.67	2.86	-	-	-	+
	8 AA779221	4.95	6.83	1.77	0.72	2.80	-	-	+	-
	2 AA455482	50.59	51.18	19.96	0.99	2.53	-	-	-	+
27853		57.48	74.35	20.99	0.77	2.74	-	-	-	+
)5 AA625788	2.18	2.08	0.89	1.05	2.46	-	*	-	+
25172		4.88	3.87	2.17	1.26	2.25	-	-	-	+
	92 AA669451	8.02	8.08	3.19	0.99	2.51	-	-	-	+
	73 AA046489	54.67	58.82	21.22	0.93	2.58		-	-	+
	63 AA486403	11.11	10.81	4.49	1.03	2.48	2	•	-	+
	88 AA418943	3.64	3.03	1.58	1.20	2.30		-	•	+
	48 AA489478	31.17	29.51	12.76	1.06	2.44		•	•	+
	42 AA976544	3.11	3.33	1.21	0.93	2.56	.	-	•	+
466	1	5.25	2.02	6.01	2.61	0.87	I.	+	-	-
	36 AA630376		5.26	2.37	1.08	2.40	:	-	-	+
	87 AA970720		3.27	88.0	0.74	2.73	1	-	-	+
	80 AA454634		4.22	1.15	0.74	2.72		-	•	+
	15 AA487680		5.15		2.36	1.10	5	-	-	-
1498			4.56		1.33	2.13		•	+	-
4616	699 AA682274	3.44	1.35	3.84	2.55	0.90)	-	-	~

gasgal	AA609456	<i>c</i> 40	4.46	3.21	1.44	2.00	_		-	+
	AA490456	6.42 0.78	0.80	0.32	0.98	2.46	_	_	+	•
	AA699926	10.36	11.94	4.03	0.56	2.57	-	-	+.	
84078	T70999	0.34	0.26	0.16	1.29	2.14		•	-	+
415415	W81118	9.68	10.64	3.83	0.91	2.52	-	•	•	+
285226	N66278	25.74	40.08	9.23	0.64	2.79		-	+	-
41207		2.05	2.42	0.80	0.85	2.57	-	-	-	+
122239	R56771	6.50	5.28	2.98	1.23	2.18	_	_	-	+
L	T98662	2.87	2.71	1.22	1.06	2.34	_	_	_	+
284865	N66734		3.47	3.15	1.61	1.78	_	_	_	+
50562	AA780841	5.61 34.20	30.09	15.14	1.14	2.26	_	_	+	-
71672	H16903 T58002	34.20 4.77	4.16	2.13	1.15	2.23	_	_	•	+
	AA135870	31.62	13.68	30.14	2.31	1.05	-	+	-	-
	AA017544	4.53	6.49	1.70	0.70	2.66	-	-	+	-
111389	T84308	4.55 2.59	2.15	1.70	1.20	2.14	_	-	_	+
	AA416782	0.75	0.75	0.32	1.01	2.32	•		-	+
	AA677574	1.64	1.62	0.71	1.01	2.30	_	-	+	-
	AA085748	0.97	1.24	0.71	0.78	2.52	_	-	+	-
	AA088371	7.14	7.60	3.02	0.76	2.36		-	•	+
	AA188999	7.14 5.86	7.00 3.92	3.24	1.49	1.81	_	-	•	+
	AA598652	10.34	13.66	4.08	0.76	2.54	-	-	•	+
	AA457092	9.85	10.74	4.18	0.70	2.36	-	•	-	+
22600	1	0.36	0.18	0.28	1.99	1.29	+	_	_	-
343688		6.94	5.30	3.54	1.31	1.96		_	+	_
	AA682399	8.65	14.46	3.24	0.60	2.67	-	-	-	+
	AA447503	3.99	2.02	3.12	1.98	1.28	_	+		_
845780	1	5.95	4.75	2.97	1.25	2.00	-	· •	-	+
266500	4	7.14	7.39	3.12	0.97	2.29	_	-	-	+
119851	I I	6.52	3.28	5.16	1.98	1.26	_	+	-	_
	194611 1 AA455519	2.66	3.75	1.05	0.71	2.54	_	-	-	+
	3 AA432030	5.78	4.46	2.98	1.29	1.94	_	-	+	-
	6 AA206914	15.56	14.08	7.32	1.11	2.13	_	-	-	+
	6 AA279172	5.70	2.95	4.40	1.93	1.30	_	+	-	-
	6 AA782292	2.75	2.53	1.30	1.08	2.12	١.	-	+	-
	3 AA190997	11.04	13.42	4.64	0.82	2.38	_	_	+	-
	0 AA421769	15.96	26.86	6.13	0.59	2.61	_	-	+	-
	1 AA443966	6.08	6.70	2.66	0.91	2.29	١.	-	+	-
	0 AA477283	4.71	4.39	2.24	1.07	2.10		-	+	
	2 AA001614	9.94	12.03	4.27	0.83	2.33	_	-	+	-
	0 AI000294	3.48	3.17	1.70	1.10	2.05		-	-	+
	3 AA001444	2.86	1.43	2.52	2.01	1.14		-	-	-
	8 AA486370	7.45	7.83	3.42	0.95	2.18	2	-	-	+
22006	3	4.81	8.77	1.92	0.55	2.50		-	-	+
	17 AA988630	13.38	13.90	6.58	0.96	2.03		-	+	-
4128		0.72	0.52	0.45	1.38	1.59	1	-	+	-
	87 AA677706	24.08	105.26	8.92	0.23	2.70		-	+	+
1514	1	2.44	2.74	1.21	0.89	2.01	1	-	+	-
	80 AA001434	3.81	1.72	5.56	2.21	0.69		+	-	-
2898	B .	6.67	4.11	5.46	1.62	1.22		-	-	-
	84 AA418670	8.86	14.76	4.05	0.60	2.19		-	+	-
1224			1.78	1.29	1.14	1.57	1	-	+	-
	94 AA406115		0.94	1.78	1.75	0.93	•	-	-	-
	44 AA112660		10.09	3.10	0.62	2.0		-	-	+
	41 AA478298		29.84	6.23	0.41	1.9	R .	-	+	-
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	Table 1-3								
462953 AA682423	3.21	5.03	1.90	0.64	1.69	•	-	+	-
1492202 4 4875013	2.60	1 02	2.70	1.31	0.96	+	-	•	-

GenBank A	ver. of	Aver. of	Aver.	Tumor	Tumor	T/N	T/N	T/N	T/B	T/B	T/B
Clone Acc. No. 7	l umors	Normals	of BPH	/Normal	/BPH	10X, >1/5	5X,>3/5	3X,>5/5	10X,>1/5	5X,>3/5	3X,>5/5
788180 AA453562	25.18	0.81	0.50	31.24	50.69	+	+ .	+	+	+	+
646037 AA196979	24.42	1.02	0.75	23.91	32.70	+	+	+	+	+	+
1034473 AA779728	33.61	1.51	1.35	22.27	24.97	+	+		+		
281003 N50880	128.77	6.30	4.90	20.43	26.26	+	+	+	+	+	+
308682 W25202	8.99	0.87	0.27	10.37	32.91	+	+	+	+	+	+
281039 N47717	14.05	0.94	0.90	14.99	15.57	+			+		
284701 N64840	33.05	7.05	1.52	4.68	21.68				+	+	+
823940 AA490213	22.39	1.32	2.83	16.99	7.91	+	+	+	+		
42415 R60981	6.86	0.51	0.72	13.52	9.49	+	+		+		
46166 H09076	13.13	0.93	1.71	14.12	7.68	+	+		+		
784168 AA432103	53.04	7.37	3.83	7.20	13.84		+		+		+
773479 AA427899	12.32	0.65	5.99	18.85	2.05	+	+	+			
222157 H85806	8.80	1.02	0.72	8.67	12.14	+	+	+	+		+
486035 AA043109	24.75	3.47	1.86	7.13	13.32	+	+	+	+	+	+
450049 AA703396	18.37	1.83	1.80	10.05	10.21	+	+	+	+	+	+
66336 T66832	1.67	0.63	0.10	2.63	16.70				+		
35058 R45192	15.59	1.89	1.44	8.24	10.81	+	+	+	+		+
26736 R39891	9.52	3.36	0.62	2.83	15.37				+		+
593185 AA159714	28.06	5.61	2.29	5.00	12.23	+			+		+
32160 R43432	26.80	3.59	2.77	7.47	9.69	l .	+	+	+	+	+
814053 AA465495	14.66	1.70	1.75	8.61	8.38	+	+		+		
292515 N68465	16.22	2.86	1.56	5.67	10.38			+	+	+	+
730739 AA436097	36.49	7.10	3.45	5.14	10.57	+		+	+	+	+
282089 N48259	11.18	1.51	1.41	7.41	7.92	+			+		
283312 N54763	63.94	6.63	12.14	9.65	5.27	+	+	+			
291255 N72215	15.20	1.35	4.65	11.29	3.27	+	+	+			
1048810 AA621342	20.15	3.04	2.59	6.63	7.78	+	+	+	+		+
197525 H52119	9.52	1.85		5.16	9.01	1			+		+
67401 T49326	1.47	0.20		7.28	6.27	+			+		
490178 AA121266	24.55	5.63		4.36	8.99	1			+		+
1556526 AA935560	0.88	0.15		5.96	7.33	•			+		
ΙΔΕΙΝΊ ΔΑΔΙΟΛΙΚ	9.35	2.23		4.18	8.70	li .			+		
1034738 AA780190	22.40	3.90		5.74	6.82	1	+	+		+	+
259374 N31952	10.51	2.33		4.50	8.04	5	·	·	+		
757435 AA437224	365.78			6.11	6.27			+	+		+
754280 AA479284	23.83			7.46	4.85	1	+	+			
877835 AA625634	23.06			5.32	6.90	j.	+	+			+
712950 AA282272	11.53			3.59	8.58	1	•		+		
50772 H16803	18.10			2.68	9.47				+		+
1636908 AI000271	6.58			1.50	10.44				+	+	+
432194 AA679414	10.34				2.56	1	+	+	,	·	
488964 AA047260	7.77				8.08	I .	·	•	+		
1412481 AA845156	8.41				9.31				+		
283173 N45236	17.85				7.21				+		
811867 AA454963	24.57				3.30		+	+	•		
26519 R20669						3	•	•	+	+	+
687297 AA235224									,	+	+
666334 AA232249						1			+	+	+
1468260 AA884926						1			+	•	•
258300 N30680									+		
971279 AA682905							+	+	•		
1610448 AA991856							•	•	+		+

461522 A	A705237	1.21	0.65	0.14	1.87	8.74				+		
868282 A	A633957	8.99	2.01	1.47	4.48	6.12			+		+	+
280217	N47941	20.52	5.86	2.89	3.50	7.09	+			.+		
240694	H78135	4.53	0.90	0.83	5.01	5.47			+			+
321706	W35398	7.80	1.86	1.25	4.20	6.23				+		
770066	AA430545	11.43	2.60	1.93	4.40	5.94	+			+		
811603	AI732784	19.62	2.74	6.22	7.16	3.15		+	+			
51773	H23211	11.48	2.40	2.09	4.77	5.50			+			+
66317	T66815	5.50	1.76	0.78	3.13	7.06				+		
35626	R45292	9.42	1.09	6.51	8.67	1.45	+	+	+			
	AA682851	9.39	1.62	2.25	5.80	4.17	+	+				
	AA702781	14.83	7.21	1.88	2.06	7.90				+		
	AA416843	11.60	4.56	1.57	2.54	7.40				+		
	AA504783	4.27	2.51	0.52	1.70	8.20					+	+
	AA136125	35.83	12.59	5.15	2.85	6.96				+		
	AA481464	47.06	7.13	14.79	6.60	3.18		+	+			
293635	N63807	26.47	3.44	12.94	7.69	2.05	+	+	+			
	AA426110	0.86	0.37	0.12	2.32	7.41						+
	AA173189	3.58	2.33	0.44	1.54	8.16					+	+
151055	H02230	27.32	7.51	4.62	3.64	5.92				+		
	AA457172	11.15	1.82	3.27	6.12	3.41		+	+			
	AA677629	0.87	0.46	0.11	1.87	7.58				+	+	
	AA173598	3.07	2.45	0.38	1.25	8.17				+	+	+
	AI025015	5.68	0.93	1.77	6.13	3.21		+	+			
	AA129551	12.35	3.96	1.99	3.12	6.21				+		
293651	N69653	13.35	1.84	6.56	7.26	2.04	+	+	+			
	AA625666	2.15	0.29	1.14	7.39	1.88		+	+			
4	AA404486	113.14	18.10	38.24	6.25	2.96		+	+			
262053		10.75	5.15	1.52	2.09	7.08						+
81331		2.84	0.78	0.51	3.63	5.53	+			+		
in telep	AA482340	34.81	11.87	5.66	2.93	6.15					+	
	AA488732	16.60	5.37	2.78	3.09	5.98				+		
	AA459761	34.94	17.03	5.05	2.05	6.92				+		+
359661	AA011095	26.24	8.79	4.39	2.99	5.97						+
		10.75	2.29	2.53	4.69	4.25			+			+
50188 = 84955		5.26	1.01	1.43	5.21	3.69		+				
159608	H16152	14.43	4.14	2.69	3.49	5.36				+		
1390582	AA843592	6.54	1.91	1.21	3.42	5.41				+		
	6 AA504842	0.92	0.60	0.13	1.54	7.26					+	
2216		20.24	13.40	2.80	1.51	7.24					+	+
	B AA452278	5.09	1.27	1.08	4.01	4.71					+	
	3 AA442991	49.03	6.94	29.52	7.06	1.66	+	+	+			
	9 AA521327	0.75	0.54	0.10	1.38	7.27				+		+
	6 AA630006	33.46	13.42	5.43	2.49	6.16				+		
	2 AA491370	14.07	9.48	1.97	1.48	7.16				+		+
	8 AA070495	21.14	3.89	6.62	5.44	3.20	Ì	+	+			
	2 AA465338	0.81	0.47	0.12	1.72	6.83				+	+	
	3 AA521416	1.31	0.47	0.12	1.61	6.93				•	+	
	O AA779919	8.90	1.72	2.71	5.19	3.28			+			
41570		3.36	0.61	1.15	5.53	2.93		+	,			
	0 W 83097 3 AA481144		0.61	0.19	2.09	6.36		•			+	
	58 AA464568		3.00	4.01	4.83	3.61	1		+		-	
	86 AA464366 20 AA464689		0.39	0.77	5.60	2.82	1	+	•			
	31 AA010932		2.50	1.49	3.13	5.26		•				+
22710	, AAVIVYJZ	7.02	2.50	1.47	٠.١٦	3.20	ŧ					•

795265 AA451807	4.58	1.26	0.97	3.63	4.74				+		
840306 AA485333		1.26	0.39	1.97	6.38					+	
824723 AA488986		4.37	2.84	3.23	4.96						+
186918 H43317		1.10	0.56	2.76	5.38				+		
191904 H38880		1.44	2.75	5.32	2.79		+	+			
771317 AA476234	9.44	1.78	3.46	5.31	2.73			+			
826072 AA521394		0.83	0.18	1.44	6.59				+	+	
30428 R42061		0.99	1.16	4.35	3.68	+			+		
1056172 AA620995	10.33	4.78	1.77	2.16	5.84				+		
251806 H96647	2.94	1.83	0.46	1.60	6.39						+
305538 N89861	11.44	3.31	2.54	3.46	4.50					+	
52933 H29315	3.59	1.99	0.58	1.80	6.14						+
824510 AA490522	1.38	1.09	0.21	1.27	6.65					+	
859228 AA666366	85.06	37.74	15.20	2.25	5.60						+
362755 AA018675	5 16.82	3.11	6.91	5.41	2.43			+			
253132 H89036	5 1.40	1.21	0.21	1.16	6.66						+
233419 H77706	5.55	1.14	1.90	4.87	2.93		+				
1592006 AA953229	9 5.30	1.51	1.24	3.52	4.28			+			+
950778 AA60863	6 3.00	1.87	0.48	1.60	6.18				+		+
951313 AA62059	1 6.84	6.68	1.01	1.03	6.75				+		
771048 AA42739	5 10.54	2.90	2.58	3.63	4.08						+
840944 AA48653	3 79.86	42.51	13.69	1.88	5.83				+		
_268211 N3632	7 3.43	0.70	1.23	4.93	2.78	+					
840606 AA48791	4 15.85	7.18	2.88	2.21	5.50						+
796161 AA46109	2 30.31	5.27	15.52	5.75	1.95		+	+			
811582 AA45851	6 10.13	5.34	1.75	1.90	5.78						+
449112 AA77748		7.72	19.71	5.51	2.16		+	+			
150262 H0112		3.31	3.87	4.12	3.53			+			
788655 AA44983		2.78	1.58	2.76	4.86						+
509731 AA04569		23.94	6.63	1.65	5.95						+
565863 AA13638		2.97	0.63	1.32	6.27				+		
509887 AA05470		7.39	17.54	5.34	2.25		+	+			
796646 AA4614	67 6.21	4.35	1.01	1.43	6.16				+		+
814235 AA4655		6.41	2.19	1.93	5.65					+	
53022 R158		0.70	0.24	1.90	5.63						+
= 377246 AA0553	06 12.60	3.02	3.78	4.17	3.34	+					
= 814662 AA4810	45 1.12	0.67	0.19	1.67	5.84					+	
51700 H228		1.21	0.82	3.02	4.47						+
80050 T638		3.84	1.52	2.12	5.35				+		
594994 AA1723		10.98	2.38	1.33	6.14						+
795907 AA4603		1.25	1.71	4.31	3.15	+					
824126 AA4906		0.37	0.10	1.58	5.87					+	
511952 AA1006		92.92	45.17	2.43	5.00	1			+		
823615 AA4969		3.54	1.97	2.65	4.76				+		
784130 AA4320		6.93	3.50	2.49	4.93						+
1535596 AA9183		285.67	66.91	1.41	6.00						+
1605407 AA988		4.61	1.65	1.95	5.45	1			+		
1606837 AA996		9.09	4.98	2.61	4.77	I .				+	
796239 AA460		23.22	4.18	1.12	6.24	1			+	+	
49302 H15			2.90	5.68	1.68						
754358 AA436			2.26	1.71	5.64					+	
207029 R98			45.41	5.54	1.80		+	+			
838285 AA458				0.99		1					+
1636108 AI015				2.47							+

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771053 AA430718 16.14 6.63 3.34 2.43 4.83 + + + 289071 NG3609 37.48 15.32 7.78 2.45 4.82 + + 289071 NG3609 37.48 15.32 7.78 2.45 4.82 + + 263697 149672 4.23 0.93 1.58 4.57 2.68 + + 263697 149672 4.23 0.93 1.58 4.57 2.68 + + + + + + + + + + + + + + + + + + +
1889071 No.
1637282
263697 H996T2 4.23 0.93 1.58 4.57 2.68
S86780 AA130677
##
839968 A1791173 12.10 2.86 4.21 4.23 2.87 287728 N75893 2.76 0.70 0.89 3.97 3.10 + 842927 AA486445 4.24 3.59 0.72 1.18 5.88 878281 AA670286 6.80 3.30 1.37 2.06 4.97
287728 N75893
842927 A4486445
876281 AA670286 6.80 3.30 1.37 2.06 4.97 359701 AA011176 5.21 2.32 1.09 2.25 4.78 897807 AA598631 5.08 2.77 0.98 1.83 5.16 855749 AA663983 50.50 12.03 18.24 4.20 2.77 264895 N21170 7.86 1.63 3.70 4.82 2.12 + 825207 AA504120 11.96 2.92 4.19 4.09 2.86 461476 AA705041 0.78 0.61 0.14 1.28 5.64 449275 AA777705 3.81 0.83 1.64 4.58 2.32 + 755239 AA422139 17.93 9.29 3.62 1.93 4.95 259973 N47829 1.66 1.95 0.28 0.85 5.99 290370 N62301 0.62 0.48 0.11 1.30 5.53 4773287 AA425534 15.94 8.17 3.27 1.95 4.87 4773287 AA425534 15.94 8.17 3.27 1.95 4.87 4773287 AA677165 20.56 33.95 3.34 0.61 6.16 4545317 R62352 7.72 6.17 1.42 1.25 5.44 462097 AA705376 8.32 6.33 1.55 1.31 1.81 4.92 462097 AA705376 8.32 6.33 1.55 1.31 5.36 + 462097 AA705376 8.32 6.33 1.55 1.31 5.36
359701 AA011176 5.21 2.32 1.09 2.25 4.78 897807 AA598531 5.08 2.77 0.98 1.83 5.16 855749 AA663983 50.50 12.03 18.24 4.20 2.77 264895 N21170 7.86 1.63 3.70 4.82 2.12 + 825207 AA504120 11.96 2.92 4.19 4.09 2.86 461476 AA705041 0.78 0.61 0.14 1.28 5.64 449275 AA777705 3.81 0.83 1.64 4.58 2.32 + 755239 AA422139 17.93 9.29 3.62 1.93 4.95 259973 N47829 1.66 1.95 0.28 0.85 5.99 290370 N62301 0.62 0.48 0.11 1.30 5.53 37173287 AA422534 15.94 8.17 3.27 1.95 4.87 3717287 AA422534 15.94 8.17 3.27 1.95 4.87 3717289 AA873159 5.15 1.21 2.03 4.26 2.53 + 3717289 AA873159 5.15 1.21 2.03 4.26 2.53 + 3717289 AA77165 20.56 33.95 3.34 0.61 6.16 + 371828549 AA976714 6.44 3.55 1.31 1.81 4.92 + 3718287 AA6977165 20.56 33.95 3.34 0.61 6.16 + 3718287 AA91761 AA677165 20.56 33.95 3.34 0.61 6.16 + 3718287 AA91761 AA697069 4.64 3.55 1.31 1.81 4.92 + 3718287 AA90609 4.64 2.75 0.94 1.69 4.95 + 371828 AA699323 38.01 9.06 15.59 4.20 2.44 + 371828 AA699323 38.01 9.06 10.68 3.77 1.71 4.85 + 371815 T61271 14.99 17.48 2.63 0.86 5.70 + 371816 AA159100 18.28 10.68 3.77 1.71 4.85 + 371828 AA699324 0.88 0.86 0.16 1.03 5.52 +
897807 AA598531 5.08 2.77 0.98 1.83 5.16
855749 AA663983 50.50 12.03 18.24 4.20 2.77 264895 N21170 7.86 1.63 3.70 4.82 2.12 825207 AA504120 11.96 2.92 4.19 4.09 2.86 461476 AA705041 0.78 0.61 0.14 1.28 5.64 449275 AA777705 3.81 0.83 1.64 4.58 2.32 + 755239 AA422139 17.93 9.29 3.62 1.93 4.95 259973 N47829 1.66 1.95 0.28 0.85 5.99 290370 N62301 0.62 0.48 0.11 1.30 5.53 279328 AA42534 15.94 8.17 3.27 1.95 4.87 472689 AA873159 5.15 1.21 2.03 4.26 2.53 + 47472689 AA873159 5.15 1.21 2.03 4.26 2.53 + 4817472689 AA877165 20.56 33.95 3.34 0.61 6.16 + 481585549 AA976714 6.44 3.55 1.31 1.81 4.92 + 491764 AA150502 0.81 0.59 0.15 1.36 5.37 + 491764 AA150502 0.81 0.59 0.15 1.36 5.37 + 462097 AA705376 8.32 6.33 1.55 1.31 5.38 + 47402097 AA705376 8.32 6.33 1.55 1.31 5.38 + 4824122 AA490609 4.64 2.75 0.94 1.69 4.95 + 4824122 AA490609 1.828 10.68 3.77 1.71 4.85 + 4814306 AA459100 18.28 10.68 3.77 1.71 4.85 + 4814306
264895 N21170 7.86 1.63 3.70 4.82 2.12
825207 AA504120 11.96 2.92 4.19 4.09 2.86 461476 AA705041 0.78 0.61 0.14 1.28 5.64 449275 AA777705 3.81 0.83 1.64 4.58 2.32 755239 AA422139 17.93 9.29 3.62 1.93 4.95 259973 N47829 1.66 1.95 0.28 0.85 5.99 290370 N62301 0.62 0.48 0.11 1.30 5.53 2773287 AA425534 15.94 8.17 3.27 1.95 4.87 4773287 AA425534 15.94 8.17 3.27 1.95 4.87 45472689 AA873159 5.15 1.21 2.03 4.26 2.53 + 45472689 AA873159 5.15 1.21 2.03 4.26 2.53 + 4548317 AA677165 20.56 33.95 3.34 0.61 6.16 + 4548317 AA677165 20.56 33.95 3.34 0.61 6.16 + 45491764 AA150502 0.81 0.59 0.15 1.31 1.81 4.92 + 462097 AA705376 8.32 6.33 1.55 1.31 5.38 + 479482 N48804 8.66 3.25 2.16 2.67 4.00 + 884122 AA490609 4.64 2.75 0.94 1.69 4.95 + 896914 AA779391 18.03 7.40 4.35 2.44 4.14 + 884306 AA459100 18.28 10.68 3.77 1.71 4.85 + 896914 AA779391 18.03 7.40 4.35 2.44 4.14 + 884306 AA459100 18.28 10.68 3.77 1.71 4.85 + 77915 T61271 14.99 17.48 2.63 0.86 5.70 + 460116 N32044 0.88 0.86 0.16 1.03 5.52 +
A61476 AA705041 0.78 0.61 0.14 1.28 5.64
449275 AA777705 3.81 0.83 1.64 4.58 2.32 + 755239 AA422139 17.93 9.29 3.62 1.93 4.95 + + 259973 N47829 1.66 1.95 0.28 0.85 5.99 + + + 290370 N62301 0.62 0.48 0.11 1.30 5.53 + <td< td=""></td<>
755239 AA422139 17.93 9.29 3.62 1.93 4.95 259973 N47829 1.66 1.95 0.28 0.85 5.99 290370 N62301 0.62 0.48 0.11 1.30 5.53 773287 AA425534 15.94 8.17 3.27 1.95 4.87 472689 AA873159 5.15 1.21 2.03 4.26 2.53 + 252314 H87175 0.69 0.62 0.12 1.11 5.66 454317 AA677165 20.56 33.95 3.34 0.61 6.16 454317 AA677165 20.56 33.95 3.34 0.61 6.16 454317 AA677165 20.56 33.95 3.34 0.61 6.16 491764 AA150502 0.81 0.59 0.15 1.36 5.37 491764 AA150502 0.81 0.59 0.15 1.36 5.37 462097 AA705376 8.32 6.33 1.55 1.31 5.38 279482 N48804 8.66 3.25 2.16 2.67 4.00 824122 AA490609 4.64 2.75 0.94 1.69 4.95 1031583 AA609323 38.01 9.06 15.59 4.20 2.44 + 824122 AA490609 4.64 2.75 0.94 4.10 3.81 2.81 1381306 AA459100 18.28 10.68 3.77 1.71 4.85 814306 AA459100 18.28 10.68 3.77 1.71 4.85 77915 T61271 14.99 17.48 2.63 0.86 5.70 260116 N32044 0.88 0.86 0.16 1.03 5.52
259973 N47829 1.66 1.95 0.28 0.85 5.99 290370 N62301 0.62 0.48 0.11 1.30 5.53 773287 AA425534 15.94 8.17 3.27 1.95 4.87 472689 AA873159 5.15 1.21 2.03 4.26 2.53 252314 H87175 0.69 0.62 0.12 1.11 5.66 454317 AA677165 20.56 33.95 3.34 0.61 6.16 454317 AA677165 20.56 33.95 1.31 1.81 4.92 491764 AA150502 0.81 0.59 0.15 1.36 5.37 491764 AA150502 0.81 0.59 0.15 1.36 5.37 462097 AA705376 8.32 6.33 1.55 1.31 5.38 279482 N48804 8.66 3.25 2.16 2.67 4.00 279482 N48804 8.66 3.25 2.16 2.67 4.00 2824122 AA490609 4.64 2.75 0.94 1.69 4.95 1031583 AA609323 38.01 9.06 15.59 4.20 2.44 896914 AA779391 18.03 7.40 4.35 2.44 4.14 896914 AA779391 18.03 7.40 4.35 2.44 4.14 814306 AA459100 18.28 10.68 3.77 1.71 4.85 77915 T61271 14.99 17.48 2.63 0.86 5.70 260116 N32044 0.88 0.86 0.16 1.03 5.52
290370 N62301 0.62 0.48 0.11 1.30 5.53 773287 AA425534 15.94 8.17 3.27 1.95 4.87 1472689 AA873159 5.15 1.21 2.03 4.26 2.53 252314 H87175 0.69 0.62 0.12 1.11 5.66 454317 AA677165 20.56 33.95 3.34 0.61 6.16 4545317 AA677165 20.56 33.95 1.31 1.81 4.92 491764 AA150502 0.81 0.59 0.15 1.36 5.37 462097 AA705376 8.32 6.33 1.55 1.31 5.38 279482 N48804 8.66 3.25 2.16 2.67 4.00 824122 AA490609 4.64 2.75 0.94 1.69 4.95 250869 N23605 123.92 32.54 44.10 3.81 2.81 896914 AA779391 18.03 7.40 4.35 2.44 4.14 814306 AA459100 18.28 10.68 3.77 1.71 4.85 77915 T61271 14.99 17.48 2.63 0.86 5.70 260116 N32044 0.88 0.86 0.16 1.03 5.52
773287 AA425534 15.94 8.17 3.27 1.95 4.87 1472689 AA873159 5.15 1.21 2.03 4.26 2.53 252314 H87175 0.69 0.62 0.12 1.11 5.66 454317 AA677165 20.56 33.95 3.34 0.61 6.16 451585549 AA976714 6.44 3.55 1.31 1.81 4.92 491764 AA150502 0.81 0.59 0.15 1.36 5.37 462097 AA705376 8.32 6.33 1.55 1.31 5.38 279482 N48804 8.66 3.25 2.16 2.67 4.00 824122 AA490609 4.64 2.75 0.94 1.69 4.95 250869 N23605 123.92 32.54 44.10 3.81 2.81 896914 AA779391 18.03 7.40 4.35 2.44 4.14 814306 AA459100 18.28 10.68 3.77 1.71 4.85 77915 T61271 14.99 17.48 2.63 0.86 5.70 260116 N32044 0.88 0.86 0.16 1.03 5.52
1472689 AA873159 5.15 1.21 2.03 4.26 2.53 252314 H87175 0.69 0.62 0.12 1.11 5.66 454317 AA677165 20.56 33.95 3.34 0.61 6.16 15885549 AA976714 6.44 3.55 1.31 1.81 4.92 491764 AA150502 0.81 0.59 0.15 1.36 5.37 139771 R62352 7.72 6.17 1.42 1.25 5.44 462097 AA705376 8.32 6.33 1.55 1.31 5.38 279482 N48804 8.66 3.25 2.16 2.67 4.00 824122 AA490609 4.64 2.75 0.94 1.69 4.95 1031583 AA609323 38.01 9.06 15.59 4.20 2.44 + 896914 AA779391 18.03 7.40 4.35 2.44 4.14 896914 AA779391 18.03 7.40 4.35 2.44 4.14 814306 AA459100 18.28 10.68 3.77 1.71 4.85 77915 T61271 14.99 17.48 2.63 0.86 5.70 260116 N32044 0.88 0.86 0.16 1.03 5.52
252314 H87175 0.69 0.62 0.12 1.11 5.66 454317 AA677165 20.56 33.95 3.34 0.61 6.16 4585549 AA976714 6.44 3.55 1.31 1.81 4.92 491764 AA150502 0.81 0.59 0.15 1.36 5.37 139771 R62352 7.72 6.17 1.42 1.25 5.44 462097 AA705376 8.32 6.33 1.55 1.31 5.38 279482 N48804 8.66 3.25 2.16 2.67 4.00 824122 AA490609 4.64 2.75 0.94 1.69 4.95 824122 AA490609 4.64 2.75 0.94 1.69 4.95 1031583 AA609323 38.01 9.06 15.59 4.20 2.44 896914 AA779391 18.03 7.40 4.35 2.44 4.14 896914 AA779391 18.03 7.40 4.35 2.44 4.14 814306 AA459100 18.28 10.68 3.77 1.71 4.85 77915 T61271 14.99 17.48 2.63 0.86 5.70 260116 N32044 0.88 0.86 0.16 1.03 5.52
454317 AA677165 20.56 33.95 3.34 0.61 6.16 ## 454317 AA677165 20.56 33.95 3.34 0.61 6.16 ## 491764 AA150502 0.81 0.59 0.15 1.36 5.37 ## 139771 R62352 7.72 6.17 1.42 1.25 5.44 ## 462097 AA705376 8.32 6.33 1.55 1.31 5.38 ## 279482 N48804 8.66 3.25 2.16 2.67 4.00 ## 824122 AA490609 4.64 2.75 0.94 1.69 4.95 ## 1031583 AA609323 38.01 9.06 15.59 4.20 2.44 ## 1031583 AA609323 38.01 9.06 15.59 4.20 2.44 ## 1896914 AA779391 18.03 7.40 4.35 2.44 4.14 ## 1896914 AA779391 18.03 7.40 4.35 2.44 ## 1896914 AA779391 18.03 18.0
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491764 AA150502
139771 R62352 7.72 6.17 1.42 1.25 5.44 462097 AA705376 8.32 6.33 1.55 1.31 5.38 279482 N48804 8.66 3.25 2.16 2.67 4.00 824122 AA490609 4.64 2.75 0.94 1.69 4.95 1031583 AA609323 38.01 9.06 15.59 4.20 2.44 250869 N23605 123.92 32.54 44.10 3.81 2.81 896914 AA779391 18.03 7.40 4.35 2.44 4.14 814306 AA459100 18.28 10.68 3.77 1.71 4.85 77915 T61271 14.99 17.48 2.63 0.86 5.70 260116 N32044 0.88 0.86 0.16 1.03 5.52
139771 R02332 7.72 0.17 1.42 1.25 3.44 4.62097 AA705376 8.32 6.33 1.55 1.31 5.38 279482 N48804 8.66 3.25 2.16 2.67 4.00
279482 N48804 8.66 3.25 2.16 2.67 4.00 824122 AA490609 4.64 2.75 0.94 1.69 4.95 1031583 AA609323 38.01 9.06 15.59 4.20 2.44 + 250869 N23605 123.92 32.54 44.10 3.81 2.81 896914 AA779391 18.03 7.40 4.35 2.44 4.14 814306 AA459100 18.28 10.68 3.77 1.71 4.85 77915 T61271 14.99 17.48 2.63 0.86 5.70 260116 N32044 0.88 0.86 0.16 1.03 5.52 +
824122 AA490609
1031583 AA609323 38.01 9.06 15.59 4.20 2.44 +
250869 N23605 123.92 32.54 44.10 3.81 2.81 +
814306 AA459100 18.28 10.68 3.77 1.71 4.85 77915 T61271 14.99 17.48 2.63 0.86 5.70 260116 N32044 0.88 0.86 0.16 1.03 5.52
*** 814306 AA459100 18.28 10.68 3.77 1.71 4.85 *** 77915 T61271 14.99 17.48 2.63 0.86 5.70 *** 260116 N32044 0.88 0.86 0.16 1.03 5.52 *** *** *** *** *** *** *** **
77915 T61271 14.99 17.48 2.63 0.86 5.70 + 260116 N32044 0.88 0.86 0.16 1.03 5.52 +
260116 N32044 0.88 0.86 0.16 1.03 5.52 +
210703 1103373 37.02 7.07 1817
703943 AA278473 4.77 2.81 0.99 1.70 4.82 +
824267 AA491265 0.74 0.41 0.16 1.81 4.68 +
813973 AA455689 3.24 0.88 1.18 3.70 2.75 +
825297 AA504536
838807 AA457681 1.90 2.58 0.33 0.74 5.71 +
1607765 AA989473 3.96 1.73 0.96 2.29 4.14
86160 T72336 2.10 0.74 0.59 2.82 3.58
1456160 AA862465 41.61 40.32 7.76 1.03 5.36 +
897767 AA598470 2.07 0.49 0.98 4.25 2.12 +
1466631 AA883679 1.21 0.71 0.26 1.71 4.63
280763 N50563 0.53 0.50 0.10 1.05 5.28 +
768643 AA430351 41.57 23.92 9.10 1.74 4.57 +
610124 AA171459 1.70 1.73 0.32 0.98 5.32 +
1631849 AI004315 25.01 12.24 5.91 2.04 4.24 +
204111 H55907 3.38 0.87 1.40 3.86 2.42 +

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491184	AA137144	6.35	1.83	2.27	3.47	2.80	+					
824237	AA491249	74.89		19.54	2.38	3.83						+
24208	R38412	7.49	3.51	1.85	2.13	4.05						+
267419		0.97	0.96	0.19	1.01	5.16						+
	AA412250	0.53	0.58	0.10	0.92	5.24				•		+
	AA680244	51.58		10.98	1.44	4.70				+		
124252		6.06	4.74	1.25	1.28	4.85				+		+
	AA160080	1.08	1.02	0.21	1.07	5.04						T
134969		6.81	4.59	1.48	1.48	4.58				+		
	AA680186	5.55	1.53	2.30	3.62	2.41	+					+
	AA459765	2.34	1.80	0.50	1.30	4.72			,			т
	AA083485	12.81	2.43	17.02	5.26	0.75		+	+		+	
	AA489696	0.49	0.34	0.11	1.42	4.58 4.10					т	+
	AA679569	4.23	2.23	1.03	1.89	4.10					+	•
	AA083671	8.46	7.83	1.73	1.08 1.10	4.86					+	
	AA425773	21.90	19.94	4.51 0.20	0.76	5.19					•	+
	AA405597	1.03	1.35	3.58	2.33	3.58						+
	AA864875	12.82	5.49 0.23	0.11	1.89	4.00						+
50764		0.43	1.65	1.02	2.24	3.62				+		
259462		3.69 71.61	18.49	36.97	3.87	1.94	+			•		
	5 AA029490	0.85	1.00	0.17	0.85	4.96	•					+
10000	6 AA421256		6.61	1.91	1.30	4.50				+		
197779	9 R93515 5 AA902831 8 AA126009	8.61 0.85	0.55	0.20	1.55	4.25						+
51142	8 AA126009	24.31	24.09	5.08	1.01	4.79						+
75770	6 AA443969	5.25	4.09	1.17	1.28	4.47						+
26446		0.77	0.84	0.16	0.91	4.79						+
5 2 2	2 AA176607	0.87	0.87	0.19	0.99	4.63						+
163734	3 AI015359	5.03	4.49	1.12	1.12	4.50						+
18676		9.43	5.75	2.39	1.64	3.95				+		
T., 2	37 AA181767	1.07	0.78	0.25	1.37	4.22						+
	2 AA487213	7.41	3.64	2.09	2.04	3.54				+		
25151		0.73	0.80	0.16	0.91	4.64						+
2 4	24 AA664101	17.75	8.17	5.31	2.17	3.34				+		
84122	20 AA486723	1.46	1.23	0.34	1.18	4.32						+
163119	94 AA994757	21.83	14.39	5.47	1.52	3.99					+	
2514		0.73	0.90	0.16	0.81	4.68						+
7674	06 AA417911	0.44	0.45	0.10	0.98	4.45						+
7452	83 AA625567	7.94	5.76	1.97	1.38	4.02						+
8150	17 AA465258	14.81	9.71	3.84	1.52	3.85						+
3468	89 W79834	0.89	0.21	0.79	4.21	1.12		+				
8238	95 AA490490	1.57	1.72	0.36	0.91	4.40						+
15533	06 AA934762		5.52	2.06	1.43	3.84						+
	22 AA001403		5.49	2.55	1.67	3.59						+
2903			17.02	4.07	1.00	4.20						++
	519 AA988798		117.37	33.70	1.15	4.01						Ŧ
	300 AA485216		1.44		0.84	4.25					+	
	672 AA056001		1.39		1.87	3.20				+		+
2560			1.47		1.17	3.88						+
	578 AA49711				1.06							+
	152 AA77543				1.20							+
	922 AA26255				0.97							•
	525 AA45211				,							+
1594	019 AA98870	1 24.27	24.53	6.55	0.99	3.70	,					•

282283 N52675	0.84	1.00	0.22	0.84	3.85
813271 AA455938	0.96	1.12	0.25	0.85	3.79
461336 AA699770	3.42	4.00	0.91	0.85	3.74
742581 AA400247	1.20	1.08	0.36	1.11	3.31
161456 H25546	13.35	10.78	5.08	1.24	2.63

Table 1-5

	Clone	GenBank Acc No.	Aver. of	Aver. of BPH	Aver.	Prostate Spec Exp	T/N & T/BPH	Secreted
-	282089	N48259	1.51	1.41	11.18	+	+	-
		AA196979	1.02	0.75	24.42	+	+	+
	745283	AA625567	5.76	1.97	7.94	+		-
	30428	R42061	0.99	1.16	4.29		+	+
	35058	R45192	1.89	1.44	15.59		+	+
	46166	H09076	0.93	1.71	13.13		+	+
	67401	T49326	0.20	0.23	1.47		+	+
	240694	H78135	0.90	0.83	4.53		+	+
	280217	N47941	5.86	2.89	20.52		+	+
	292515	N68465	2.86	1.56	16.22		+	+
	593185	AA159714	5.61	2.29	28.06		+	+
	730739	AA436097	7.10	3.45	36.49		+	+
	1592006	AA953229	1.51	1.24	5.30		+	+
	32160	R43432	3.59	2.77	26.80		+	
	42415	R60981	0.51	0.72	6.86		+	
	50188	H17943	2.29	2.53	10.75		+	
	51773	H23211	2.40	2.09	11.48		+	
	81331	T60075	0.78	0.51	2.84		+	
	222157	H85806	1.02	0.72	8.80		+	
	281003	N50880	6.30	4.90	128.77		+	
1	281039	N47717	0.94	0.90	14.05		+	
į.	283173	N45236	3.97	2.47	17.85		+	
	308682	W25202	0.87	0.27	8.99		+	
	450049		1.83	1.80	18.37		+	
	486035		3.47	1.86	24.75	1	+	
1000	488964		2.13	0.96	7.77		+	
1	490178		5.63	2.73	24.55		+	
num.	712950	8	3.21	1.34	11.53		+	
	757435		59.86		365.78	1	+	
	784168	1	7.37		53.04		+	
	788180		0.81	0.50	25.18	1	+	
	814053		1.70 1.32		14.66 22.39	1	+	
	823940 868282		2.01		8.99		+	
ä	877835	1	4.34		23.06	4	+	
	1034473		1.51		33.61		+	
	1034738	1	3.90		22.40	1	+	
	1048810		3.04		20.15	1	+	
	141248	.	3.50		8.41	I I	+	
	155652	i .	0.15		0.88	4	+	
	2651	P.			36.05	1		+
	3562				9.42			+
	5170	l				3		+
	5293							+
	7791	B C						+
	8005							+
	8495	3						+
	12425	\$				6		+
	13496	i9 R31789	4.5	9 1.48	6.8	1		+

Table 1-5

1	GenBank	Aver.	Aver.	Aver.	Prostate	T/N	
Clone	Acc No.	of	of BPH	of	Spec Exp	& T/BPH	Secreted
150262	H01129	3.31	3.87	13.66			+
151055	H02230	7.51	4.62	27.32			+
159608	H16152	4.14	2.69	14.43			+
186768	H51914	5.75	2.39	9.43			+
186918	H43317	1.10	0.56	3.04			+
233419	H77706	1.14	1.90	5.55			+
251806	H96647	1.83	0.46	2.94			+
252314	H87175	0.62	0.12	0.69			+
256680	H96392	1.47	0.44	1.72			+
259462	N29545	1.65	1.02	3.69			+
260116	N32044	0.86	0.16	0.88			+
263697	H99672	0.93	1.58	4.23			+
280763	N50563	0.50	0.10	0.53			+
283312	N54763	6.63	12.14	63.94			+
284701	N64840	7.05	1.52	33.05			+
290370	N62301	0.48	0.11	0.62			+
290378	N64508	17.02	4.07	17.10			+
291255	N72215	1.35	4.65	15.20			+
293635	N63807	3.44	12.94	26.47			+
293651	N69653	1.84	6.56	13.35			+
321706	W35398	1.86	1.25	7.80			+
346889	W79834	0.21	0.79	0.89			+
361922	AA001403	5.49	2.55	9.18			+
375682	AA032221	12.66	6.60	31.04			+
377672	AA056001	1.39	0.81	2.59			+
416374	W86202	2.66	0.81	4.49			+
430465	AA680186	1.53	2.30	5.55			+
450307	AA682851	1.62	2.25	9.39			+
454317	AA677165	33.95	3.34	20.56			+
461336	AA699770	4.00	0.91	3.42			+
461476	AA705041	0.61	0.14	0.78			+
491184	AA137144	1.83	2.27	6.35			+
491764	AA150502	0.59	0.15	0.81			+
509731	AA045699	23.94	6.63	39.41	1		+
509887		7.39	17.54	39.47			+
511428		24.09		24.31			+
511952	AA100674	92.92		225.66			+
530185 564801	AA083671	7.83		8.46	3		+
586780	AA129551 AA130677	3.96 1.75		12.35	1		+
593658		1.73		1.92	1		+
594994		10.98		1.08 14.61	4		+
595695	•	2.33		3.58			+
609052	1	0.87		0.87	1		+
610124		1.73		1.70			+
666334	•	1.73		1.64	L .		+
731410	1	0.58		0.53	•		+
756600	1	7.13		47.06			+
757392		0.37		0.86	1		+
					1		

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Table 1-5

1	GenBank	Aver.	Aver.	Aver.	Prostate	T/N	
Clone	Acc No.	of	of BPH	10	Spec Exp	& T/BPH	Secreted
767406	AA417911	0.45	0.10	0.44			+
768643	AA430351	23.92	9.10	41.57			+
770066	AA430545	2.60	1.93	11.43			+
773248	AA425773	19.94	4.51	21.90			+
773479	AA427899	0.65	5.99	12.32			+
784130	AA432075	6.93	3.50	17.23			+
787938	AA452278	1.27	1.08	5.09			+
788655	AA449837	2.78	1.58	7.66			+
795265	AA451807	1.26	0.97	4.58			+
796161	AA461092	5.27	15.52	30.31			+
796239	AA460675	23.22	4.18	26.07			+
811582	AA458516	5.34	1.75	10.13			+
811603	AI732784	2.74	6.22	19.62			+
811867	AA454963	2.97	7.43	24.57			+
815163	AA481144	0.57	0.19	1.20			+
823615	AA496957	3.54	1.97	9.40			+
823895	AA490490	1.72	0.36	1.57			+
824126	AA490611	0.37	0.10	0.59			+
825822	AA491370	9.48	1.97	14.07			+
826072	AA521394	0.83	0.18	1.20			+
826109	AA521327	0.54	0.10	0.75			+
838285	AA458747	1.36	0.21	1.34			+
838807	AA457681	2.58	0.33	1.90			+
839081	AA488732	5.37	2.78	16.60			+
840894		11.87	5.66	34.81			+
855624		8.17	5.31	17.75			+
878152	1	3.49	1.10	4.20			+
884606		13.42		33.46	i .		+
897767	*	0.49		2.07	1		+
951313	1	6.68		6.84	3		+
1031583	L.	9.06		38.01			+
1056172	1	4.78		10.33	i		+
1456160	1	40.32		41.61	*		+
1472689		1.21		5.15			+
1535596		285.67		401.67	1		+
161044	į.	57.06		135.55			+
1631849	9 AI004315	12.24	5.91	25.01	1		т

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

n a t leas	
or higher i	
Table 2-1: T3NO vs BPH: Genes Upregulated 5 fold or higher in at least 1	
Table 2-1: T3NO vs BPH: Genes Upr	
Table 2-1: T3N	

TOE									4,007	of Tumor	old/MPM125/A	t of Tumor
	IMAGE	Gen Bank	Hodony	AveTumor	Fold(AveTumor/Av Fold(MPM120/A Fold(MPM121/A	Fold(MPM120/A	Fold(MPM121/A F	old(MPM124/A F	veBPH)	veBPH)	veBPH)	Samples
	Clone ID	Accession Number			•BPH)			1 38	52.32	22.54	10.36	စ
•	788180	AA453310	1.96	133,47	67.89	149.63	165.70	1.50	31.06	53.59	35.80	9
- ‹	648037	A A 106979	2.38	114.00	47.98	65.24	24.62	16.17	A 28	41.24	19.50	9
, ,	284704	NRARAD	2.41	73.10	30.37	31.27	50.07	(a. 6)	15.57	45.42	20.45	9
, ,	450049	44703396	2.55	76.64	30.08	38.52	11.57	4 6.04 4 0.04	19.72	21.41	63.61	9
t 4	245768	NSSSB	0.0	0.21	22.49	11.34	8.80	90.0F	15.57	11.61	13.29	9
D 4	284003	N50880	24.05	423.65	17.62	22.19	20.64	0.727	10:01	5.71	7.21	ø
۰ ۲	138605	R35051	2.56	44.38	17.34	29.78	41.52	9.20 4.40	9.24	10.23	5.96	ø
- 00	133130	R28397	1.48	25.46	17.23	30.48	41.34	 	1.31	3.32	7.01	စ
, a	838668	AA457235	3.44	46.10	13.39	11.55	50.28	11 09	10.01	17.09	10.66	æ
, £	179753	H51549	0.01	0.12	11.81	9.28	11.84	5 1	6.14	4.27	15.03	မှ
: =	1030855	AA621761	1.24	12.57	10.16	7.99	15.61 6.83	6 8 8 8 8	7.12	8.55	19.24	စာ
: 2	51433	H20747	2.07	19.56	9.43	6.78	6.32	1 00	4.72	3.65	6.31	ø
<u>.</u>	306066	N91003	0.80	6.9 46.9	8.70	13.54	0r.9r	 	5.51	12.71	6.41	9
: 7	195975	R91398	0.50	3.83	7.67	5.89	6.60	- ce	7.18	5.75	8.20	စ
. 45	21831	172543	0.29	1.97	6.87	3.24	10.05	200 1	10.04	4.07	5.11	ဗ
. 4	282217	N79180	90.0	0.42	6.73	3.94	LT.0T		5.13	5.23	11.57	ထ
: 4	825859	AA504784	1.91	12.19	6.37	6.28	40.4	000	4.79	4.07	3.73	စာ
~	428816	AA004681	0.10	0.60	6.03	6.54	\$ 6 \$ 6	 2. 2.	7.48	3.67	8.48	ထ
6	195034	R88764	0.02	0.14	5.84	3.19	9.00	3.89	8.96	3.53	3.06	9
20	359701	AA011176	2.91	12.87	4.42	3.21	90.0	4.48	3.84	3.52	3.85	9
77	243641	N49548	1.55	6.34	4 .09	3.17	0.80	o so	4.47	3.77	4.01	စ
23	262035	H98688	0.22	0.88	3.95	3.77	5. c	18.85	12.53	19.49	9.01	တ
ន	287528	N62096	0.30	3.88	12.94	15.20	26.45	45.4	17.70	1.22	3.92	ĸ
7	784168	AA432103	15.97	194.72	12.19	9.62	26.10	10.24	5.27	13.26	1.75	vo
52	259374	N31952	1.29	13.52	10.47	7.91	24.35 0.06	76	3.08	33.88	4.85	ĸ
92	210820	H67712	0.19	1.82	9.61	2.75		7.87	13.10	1.22	3.27	10
22	193923	R83407	17.59	158.73	9.05	10.87	78.73	3.79	3.02	4.81	0.54	ĸ
8	37980	R61372	0.09	0.74	8.39	9.8 0.0	10.06	6.72	12.64	2.83	7.06	ĸ
58	1031599	AA609485	0.27	2.07	7.65	1 0.00 1 0.00	10.90 A 16	4.03	15.83	2.66	4.49	ທ
စ္တ	86160	172336	1.71	11.25	6.59	85.7 85.9) (8) (8)	13.77	2.12	4.36	6.18	so.
뜐	268211	N29986	0.57	3.71	0.40 0	2						

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one turnor sample or 3 fold or higher in at least 2 tumor samples.

	IMAGE	Gen Bank Accession	AveBPH	AveBPH AveTumor	Fold(AveTumor/Av Fold(MPM121/A Fold(MPM124/A Fold(MPM122/A Fold(MPM123/A Fold(MPM125/A # of Tumor Samples VeBPH) VeBPH) VeBPH) VeBPH) VeBPH) VeBPH)	old(MPM120/A	Fold(MPM121/A F	old(MPM124/A veBPH)	Fold(MPM122/A F	old(MPM123/A F veBPH)	Fold(MPM125/A veBPH)	samples
	Cione 10	Number					600	5.48	10.02	1.94	3.81	-
32	462468	AA699895	0.38	2.39	8.34	5.85	10.02	4.27	14.38	9.64	3.94	ĸ
ಜ	866276	AA663160	3.28	20.49	6.28	1.91	3.57		408	2.69	8.45	ĸ
8	811582	AA454597	3.58	21.22	5.92	9.12	84 84		5.71	11.11	7.68	40
33	428436	AA004412	0.53	3.12	5.83	1.13	4.65 0.69	5 5	5.38	6.96	2.08	ĸ
8	130138	R20813	9.0	0.20	5.62	7.23	08.9	9.30	4	3.72	11.48	ĸ
37	347586	W81432	9 .0	0.24	5.62	2.89	/8.4 /0.	<u> </u>	11.93	6.34	3.73	ĸ
ဆွ	795729	AA460291	0.02	0.11	5.58	5.69	06.4 00.0	50.0	83	9.01	4.56	ĸ
39	825785	AA505117	1.58	8.57	5.51	3.24	6.98	S 8	3.61	13.09	3.10	١co
\$	49858	H29290	0.33	1.80	5.39	2.31	3.63	0.00	4.21	8.09	9.54	ĸ
=	417075	W87823	0.83	4.38	5.30	2.79	3.52	0 6	3.36	7.56	4.13	40
2	840751	AA486071	1.73	6 0.6	5.25	3.31	10.29	0.0	90 80	2.79	3.31	ю
5	241821	H93381	1.06	5.43	5.10	3.95	6.61	29.0	780	4.62	2.43	တ
4	29063	R40970	1.68	8.52	5.08	4.26	6.83) (i.e.	7.00	8.48	5.11	10
. 5	124128	R01348	1.25	6.16	4.92	2.53	5.09	3.08	1,43	4 .02	4.20	พ
: 5	238821	H65030	4.70	23.13	4.82	2.33	3.91	3.66	2 6	3 53	2.93	w
2	134719	R28287	0.29	1.40	4.85	5.02	7.58	80.4	t to	3.82	5,12	10
. 4	287745	N82244	3.81	18.41	4.83	2.38	4.28	3.75	4 (0) 1 (0	20:0	4.56	ĸ
2 5	24444	N34483	0	900	4.58	0.79	8.80	4.19	3.78	7.0	277	• • c
? 9	*******	20110N	0.79	3.60	4.55	1.68	4.13	3.48	7.19	\$ 0 0 0	3.63	, ka
3 3	1140471	M02538	338	15.26	4.54	9.25	5.31	0.40	5.39	3.28 1.4	8 7	. 10
5 8	100010	0040000	0 13	0.59	4.48	4.82	5.19	6.45	4.67	1.7.1		. sc
70 1	8 to 20 to 3	000000000000000000000000000000000000000	2 6	2 2 2	4.37	3.68	3.50	2.94	3.83	6.40 5 :	9.00) K
3 :	07//89	7080807	20.0	0 12	4. 5.3	3.87	6.96	3.48	5.59	40:		s ka
\$ 1	20000	1971CN	3 6	000	4.23	6.18	3.28	3.22	6.00	90.0	9 6	, w
ខ្ល	265003	180120	, a	35.37	4.16	6.16	7.98	3.20	3.85	0.24	3.50) k
8 1	108040	VV/0234	948	1 88	4.03	2.15	7.06	3.45	3.09	4.6	3.5	, u
à	610001	107/0/) ¥	7 83	4.02	3.49	4.63	2.64	1	5.85	9.09) W
8	842848	AA486281		4. C	20:4	3.25	3.93	4.50	3.68	4.50	2.31	n *
20	40699	R55882	co c		2 6	4.70	6.56	3.16	3.29	3.49	49.0	D 1
8	359661	AA011096	23.32	78.7	\$ 600 d	9 6	3.36	3.88	4.66	4.89	0.89	ıo
2	126230	R06307	0.03	0.10	3.50		3.29	3.51	3.03	3.10	1.45	10
62	375682	AA032221	15.66	45.43	2.80	50.5	1.60	59.34	105.24	0.80	19.00	₹
8	1468260	AA884926	0.65	22.36	¥.5	16.69	5 4	29.56	0.88	23.62	1.26	*
2	743828	AA634379	0.58	12.90	22.20	29.78	10, 10, 10, 10, 10, 10, 10, 10, 10, 10,	3.22	1.15	2.77	3.51	4
88	1390584	AA843592	7 .	33.09	18.02	49.03	c 6. 5.	*	! ! !			

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold of higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

	IMAGE	Gen Bank	100	AveTumor	Fold(AveTumor/Av	Fold(MPM120/A	Fold(MPM121/A	Fold(MPM124/A	AveTumor/Av Fold(MPM120/A Fold(MPM121/A Fold(MPM124/A Fold(MPM122/A Fold(MPM123/A Fold(MPM126/A # of Tumor AveTumor/Av Fold(MPM120/A Fold(MPM121/A Fold(MPM124/A Fold(MPM122/A Fold(MPM126/A # of Tumor AveBPH) veBPH) veBPH) Samples	Fold(MPM123/A	Fold(MPM126/A veBPH)	# of Tumor Samples
	Clone ID	Number			•8PH)	(HIJGRA)		,	£7 03	15.40	1.72	4
;	1	702020	070	3.34	15.84	9.65	9.81	<u>.</u>	80.75	40 24	5.35	*
8	4104/	K27784	}		41 84	2.76	4.13	1.86	8.52	19:04		•
67	74494	AA625890	08.	74:77		22 49	23.74	3.29	9.26	2.32	- e: ·	• •
89	1034473	AA779728	0.13	1.36	00:01	7 70	10.45	13.83	13.97	1.00	00.	•
69	487165	AA045074	0.01	0.09	8.96	D 4.0	24.53	1.58	8.31	4 .84	2.78	4
02	1458424	AA862992	90.0	0.69	8.20	10.16	55.12	23.6	24.39	4.57	6.15	*
	360155	AA013260	0.01	0.11	8.02	10.07	5/.0	27.7	10.46	49.	2.63	4
: £	144740	876228	0.19	1.27	6.55	7.58	10.38	0.02	2 0	10.25	15.67	*
÷ ;	24046	890060	0.93	5.97	8.42	3.07	2.31	4.37	. 60.50 60.60 60.60	2.35	3.29	4
2 7	K0318K	AA165678	9.70	60.24	6.21	9.37	12.99	2.46	0.00	1.74	10.42	4
: ;	40KR172	A A R 2 0 9 9 5	11.61	70.00	6.03	99'9	5.08	T0.0T	6 CO	1.51	2.67	•
2 5	10001	A 424407	98	5.55	5.84	8.95	11.77	5.12	2.02	. 0.5 .0.5	3.19	4
٤ ۽	102303	AA600044	9 6	0.07	5.65	16.58	0.83	6.44 4	4.81	2.03	2.77	*
: ;	20112	A 4 70 52 5 B	0.34	1.88	5.48	2.60	6.47	9.51	0 4 0 0	966	2.67	*
2 1	40104	AA16250	7.5	3.80	5.23	3.93	6.17	7.31	9.02	87.7	86.	4
2 :	00/184	AA115259	2 4	, a	40.6	6.20	9.84	6.65	1.30	0.20	9 17	4
සි	787938	AA452278	0/-	5 6		5.74	6.67	0.33	1.52	\$: i	90.0	. 4
8	754034	AA479058	20.0		¥ 93	3.02	13.78	4.92	2.90	1.71	5.20 5.00 5.00 5.00 5.00 5.00 5.00 5.00	- ◄
82	970590	AA683077	9. 9.	0.0	997	166	3.59	10.99	8.52	0.64	D. 4	• •
8	68767	T53389	0.05	0.08	A .		3.11	2.03	11.89	6.25	2.97	•
\$	292515	N68465	3.53	17.34	4 .91	3.22	. 6	6.56	2.56	2.70	10.45	4
8	296210	N74392	4.0	2.17	4.90	3.19	t 65.00	4.63	7.12	1.57	11.65	*
8	40402	R52089	0.02	0.09	4.82	3.44	S 6.9	5.7.c	9.28	4.42	5.71	*
87	858153	AA633811	0.05	0.24	4.82	1.99	08.4 08.6	t (;) t	3 55	3.98	1.23	4
. 60	704697	AA282253	0.27	1.26	4.71	1.92	9.73	8.7		1.16	2.07	*
2	814308	AA459318	9.65	45.42	4.71	5.23	10.10	- ·) 1	1.30	7.69	*
8 8	781233	AA448316	0.05	0.10	4.70	0.48	4.83	9.30	- c	1.85	2.40	4
5	823794	AA490263	1.70	7.94	4.66	3.60	10.81	60.00 00.00	5 7	1.63	4.68	*
6	280170	N32072	12.85	59.29	4.61	5.52	5.50	9.33 10.00	5 6	2.38	2.87	4
1 8	449309	AA777902	2.47	11.22	4.54	4.85	3.66	3.87	4 6.0	184	3.79	*
3 2	2284KK	LIR1758	0.98	4.35	4.51	4.04	2.42	10.26	9/:-	. Ce o	3.44	*
\$ 8	20870	VANCORR	9	4.15	4.41	1.85	2.47	3.73	9.40		17.	*
2 3	#100A7	70000	0.47	2.07	4.38	5.34	6.46	6.39	1.35	3.07		•
S :	7000R1	C1800A	; ;	8 20	4.38	2.17	5.43	7.7	1.35	3.80	7 60	. •
S	420201	AAUU4020		40 45	434	3.25	3.54	4.23	10.05	2.88		- ◄
8	391984	A1003625	7 .	19.13	4.29	5.19	4.37	3.15	9.71	1.81	 	•
œ	51700	H22854	76.	34.5	!		ო					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

Fold(AveTumor/Av Fold	Gen Bank Fold(A	Fold(A	Fold(A	Fold(AveTumor/Av Fold(M	Av Fold(M	PM120/A	Fold(MPM121/A	Fold(MPM124/A	Fold(MPM122/A veBPH)	fold(MPM123/A veBPH)	Fold(MPM125/) veBPH)	A # of Tumor Samples
Accession Avebru	Accession Aveish Aveiumos	Averamos		eBPH)		VeBPH)	(H. LOBA			101	080	•
Number	Number Ar CA 60 10	07 03		4.14		5.48	5.99	2.50	3.26	4.67	28.4	• 🔻
AA479494 15.01 02.10	15.01	05:10				1.97	3.35	4.59	7.43	0.10	97	. ◄
H66840 0.75 3.08	0.75 3.08	3.08		5 7		6.87	3.20	5.81	2.50	1.75	, t	. ◄
AA443090 0.02 0.10	AA443090 0.02 0.10	0.15 80 80		60.4		2.86	6.98	3.63	4.72	4. 20 0. 00	5 6 6	• 🕶
7.01	172 7.01	7.01		4.07		3.68	2.48	4 .06	71.7	69.0	3.20	4
MB0128 13.56 54.72	MB0128 13.56 54.72	54.72		40.4		3.89	3.21	2.71	5 13	8.53	2.31	4
AA703046 1.44	1.44 5.81	5.81		4.03		1.27	3.01	3.80 8.80	2.74	2.97	3.41	4
3 AA609106 1.74 6.94	AA609106 1.74 6.94	6.94		3.99		6.11	5.05 50.5	5.08	3.69	6.17	1.75	4
AA939275 0.61 2.38	AA939275 0.61 2.38	2.38		3.91		2.08	00.6	4	1.37	4.41	3.71	→
R76247 0.43 1.67	R76247 0.43 1.67	1.67		3.89		9.05	, 5, 5, 5, 5, 5, 5, 5, 5, 5, 5, 5, 5, 5,	. 6	2.10	3.38	2.84	*
W32272 1.23 4.75	W32272 1.23 4.75	4.75		3.84		5.19	0.74	5.19	3.29	7.01	0.18	*
AA291742 0.05 0.19	AA291742 0.05 0.19	0.19		3.78		2.57	÷ ;	3.79	6.85	1.28	0.83	→
N25883 0.28 1.07	N25883 0.28 1.07	1.07		3.78		6.40	5.33 4.88	2.58	3.38	4 .60	4.50	4
N25920 1.81 6.68	N25920 1.81 6.68	6.68		3.70		2.93	4. c	3.82	2.44	4.10	5.85	4
N53378 0.93 3.42	N53378 0.93 3.42	3.42		3.66		2.34	5.5 5.0	2.55	4.74	3.85	0.49	→
AA011320 3.58 13.06	AA011320 3.58 13.06	13.06		3.65		4.67		3.36	5.17	4.30	1.00	*
AA630104 0.98 3.56	AA630104 0.98 3.56	3.58		3.65		2.92	- i	3.62	5.06	3.32	2.93	4
H82081 0.83 3.03	H82081 0.83 3.03	3.03		3.65		4.16	Z.13	3.81	5.38	2.39	1.1	4
AA427528 10.52 38.08	AA427528 10.52 38.08	38.08		3.62		3.55	7. c. a	3.11	1.20	3.29	5.98	→
AA045709 0.98 3.54	AA045709 0.98 3.54	3.54		3.60		1.73	6.43	5.03	4.80	4.57	1.28	4
T90980 11.33 40.58	T90980 11.33 40.58	40.58		3.58		1.21	. c	3 15	2.19	2.78	4.53	*
T62842 0.03 0.10	T62842 0.03 0.10	0.10		3.58		5.81	3.02	988	1.92	5.82	3.66	4
AA664104 0.96 3.38	AA684104 0.96 3.38	3.38		3.52		2.09		3.87	3.79	2.79	1.62	*
AA454988 2.96 10.39	AA454988 2.96 10.39	10.39		3.51		3.92	2 6	3.53	2.17	2.42	4.40	→
0.36 1.25	N47886 0.36 1.25	1.25		3.49		5.3	, ,	4 85	1.73	3.48	5.29	*
AA704650 9.01 31.38	AA704650 9.01 31.38	31.38		3.48		2.44	3.12	00 6	4.08	4.15	2.06	*
AA757401 2.20 7.66	AA757401 2.20 7.66	7.66		3.48		2.26	5.5	60. 6	3.23	4.16	2.95	4
R63253 1.41 4.88	R63253 1.41 4.88	4.88		3.47		1.74	6.4. r	. t.	1.59	6.09	3.28	→
N63609 19.65 68.00	N63609 19.65 68.00	68.00		3.46		3.37	90.0	1.00	4.28	4.93	1.55	*
AA677461 0.55 1.89	AA677461 0.55 1.89	1.89		3.45		1.61	6.2.4	2) · c	3.25	0.92	0.95	*
H93335 2.71 9.21	H93335 2.71 9.21	9.21		3.40		5.12	6.73	£ 6	2.28	4.65	2.52	4
H48502 0.16 0.53	H48502 0.16 0.53	0.53		3.38		3.02	4.77	200	4.60	1.60	3.25	4
AA758154 0.68 2.24	AA758154 0.68 2.24	2.24	_	3.36		1.87	79.0 80.0	4 07	1.86	₹.90	3.98	*
209381 H64085 0.97 3.25 3.35	H64095 0.97 3.25	3.25		3.35		2.04	3.20	è				
							4					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold of higher in at least one tumbrisample or 3 fold or higher in at least 2 tumor samples.

		Gen Benk				A/NOM420/A	Fold/MPM121/A	Fold(MPM124/A	Fold(MPM122/A		Fold(MPM125/A	# of Tumor
	IMAGE	Accession	AveBPH	AveTumor	Fold(AVe LumonAv	veBPH)	veBPH)	veBPH)	veBPH)	veBPH)	VeBPH)	Samples
		Number				9	3 04	86.86	3.28	1.47	3.16	· →
\$	23897	R38364	0.38	1.24	3.28	1.98	9.81	9 4	3.37	3.97	2.04	*
135	1487905	AA883504	1.18	3.84	3.27	1.87	3.82) () ()	980	4.48	1.12	*
138	1556526	AA935560	1.17	3.77	3.21	3.58	5.83 88.63	3.20	3 6	333	3.88	4
137	491418	AA115537	0.54	1.72	3.20	1.82	3.32	2.46	9.4	F 41	1.80	4
138	744001	AA629039	3.87	12.24	3.17	1.60	3.05	3.27	0.4	- 4	2	4
5 5	293110	N63848	3.38	10.70	3.17	2.38	3.94	3.44	3.86	6.99		. •
3 5	10001	4 4 4 20 6 0 2	1 + 1	3.67	3.14	3.26	2.51	3.73	2.89	3.22	3.21	• •
2	1870787	A44.29002	- 5		80 6	2.09	3.15	1.69	3.87	3.10	4.59	4
¥	454689	AA677206	70.0	2 5	9 6	4.26	3.84	4.05	3.20	1.06	1.87	~
75	384293	AA709370	0.50	1.03	60.6	9 6	76 6	3.79	3.14	3.71	1.62	*
1 3	34091	R44132	0.53	1.63	#0.50 10.50	5 6	, e	20.6	4.16	0.88	4.58	₹
<u>‡</u>	743362	AA400385	0.03	0.09	3.02	3.33	2.6	3 6	3 17	3.53	3.40	*
145	128785	R10015	4.1	3.44	3.02	2.11	2.81	00.5	- u	222	3.94	4
148	429299	AA007370	1.28	3.83	2.98	3.22	3.29	1.89	5.55	1 4 1 7	3 33	~
	136819	R36465	0.47	1.37	2.84	3.15	2.95	3.25	3.12	6.6) H	
: :	1323K)	AA400804	£2 £3	123.16	2.88	3.66	5.22	3.12	1.22	0.47	c: co	٠,
2 :	00000	£0000 / ¥~	3		27.6	3.57	4.62	3.03	1.81	0.36	3.03	• (
3	0/200/	AA480348		3 6	25.02	33.	1.13	1.13	14.34	134.44	1.13	י פא
3	/180/7	N3Z514	5 6	9 6	15.42	64.92	1.22	0.11	0.61	3.63	22.02	rs .
151	782233	AA431721	0.20	2 6	4 6 6	1 39	3.75	0.97	71.36	3.78	2.68	က
152	320495	W16659	90.0	0.86	13.88	9 - 6	1.52	4.77	26.47	44.95	2.05	က
153	283579	N94111	98:0	12.90	13.42	5 6	25.22	5.01	2.22	0:30	1.22	က
2 5	281039	N47717	0.78	8.88 8	12.85	01.5	1 1 0 0 0 TO 0	1.83	0.27	18.31	0.82	က
155	490718	AA115761	9.0	0.48	12.37	40.04	5 6	200	55.00	5.81	3.6	ო
158	796747	AA443147	0.08	0.92	11.12	0.12	21.0	7 7	1.25	0.86	1.67	60
157	287728	N59158	0.51	5.43	10.69	20.89	31.83	 		1.07	2.28	ო
158	282089	N48259	1.90	19.34	10.17	21.89	29.40		5 6	87.0	4.21	ო
159	731290	AA416627	4.08	33.97	8.32	14.84	26.99	1.81	97.1	32.28	2.68	က
8	470216	AA028987	2.85	23.62	8.02	1.97	4.75	00.1	8.5	00.76	2.42	ez
161	375827	AA039851	0.07	0.57	7.75	3.54	3.42	1.63	1.38	0. % 0. %	; ¢	• e:
162	277714	N49577	0.13	9.0	7.33	0.87	3.83	1.15	32.30	5.08		, (
1 2	757451	AA437241	0.47	3,37	7.13	4.67	30.78	1.10	3.88	0.99	وي: د	, c
3	486035	A A O 4 O B B 1	5.25	35.89	6.84	2.41	4.29	2.60	21.59	8.58	AC.	, ,
3 4	200743	174134	80	870	6.33	13.15	16.83	1.72	1.58	1.47	3.24	o (
3 4	EREC 148	4 4 2 2 2 2 0 0 0	3 2	200	6.24	1.91	7.96	4.25	20.28	2.26	0.81	m (
8 5	757389	AA437142	0.02	0.11	5.76	1.70	4.23	0.65	4.75	0.56	22.68	m
<u>:</u>))	; ;	:				5					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold of higher if at least one tumor samples.

A # of Tumor Samples	c	- o	က	က	ო	ო	ဗ	n	· e	,	n (n	က	က	က	က	က	က	က	ო	က	က	က	က	က	က	က	က	က	က	က	က	က	က	က	က	
Fold(MPM125/ veBPH)	Š	5 .7	2.26	2.12	0.10	2.62	6.86	8.28	76 4	t 0.0	3.52	4.38	0.89	2.65	0.13	3.58	7.68	2.38	0.38	1.83	1.27	8.83	9.67	4.11	3.80	1.74	6.79	9.45	8.82	1.39	1.52	2.08	5.34	4.00	2.22	11.30	
Fold(MPM123/A		4.09	1.05	0.88	17.97	0.89	11.48	7 48	P : .	1.13	2.89	7.22	3.94	1.66	4.55	2.45	1.05	0.26	2.43	46.0	1.09	1.24	790	8,96	15.21	1.01	1.92	0.68	1.37	77.	69	1.15	0.79	86 7	4 58	4	2
Fold(MPM122/A		16.65	1.32	0.72	0.13	20.36	7 6 7	6. 6	66.0	0.90	12.13	0.98	4.85	1.93	1.37	13.00	8 18	25.0	883	3	t 600	 	27.0	9.6		9.50 4.00 8.00		2 0	1.50	3. 5	90.0	338	3 2		7. P	7.80	1.80
-old(MPM124/A	Vebruj	2.16	4.47	8 83		4 : C	0.73	2.60	8.08	2.17	6.84	13.33	5		5 7		75.0	Z.00	e 6	87.7		4. 0. 4	1.80	0.5 8 8	0.20	0.68	% t	0:0	Ö. 6	2.68	7.47	0.0	4.10	0.28	7.11	97.5	3.64
Fold(MPM121/A	veBPH)	1.65	10 50	6.0	14.24	re:D	4.09	4.85	2.75	12.12	1.90	1 24	17.30	87.4	c0.8	16.84 4.3	1.49	6.21	8.45	8.33	9.86	12.69	7.96	1.67	0.16	1.96	9.76	2.40	7.99	0.81	1.98	6.18	2.50	7.63	2.29	3.44	1.55
Fold(MPM120/A	veBPH)	7,	1	5.83	6.81	6.26	3.13	2.72	2.52	7.68	1.82	4 6	1.52	1.96	5.69	0.67	1.47	1.67	7.28	1.90	8.62	4.09	0.98	3.30	10.35	0.46	7.32	7.69	5.29	3.45	3.07	2.09	10.98	0.28	2.52	2.93	3.05
Fold(AveTumor/Av Fold(MPM120/A Fold(MPM121/A Fold(MPM124/A Fold(MPM122/A Fold(MPM123/A Fold(MPM126/A # of Tumor	eBPH)		5.60	5.59	5.58	5.45	5.30	5.03	50	.00	A: +	cs.4	4.78	4.64	4.62	4.59	4.58	4.56	4.52	4.49	4.	4.40	4.34	4.33	4.33	4.31	4.28	4.28	4.16	4.12	4.06	4.03	4.02	3.84	3.92	3.90	00 6
	AveTumor		3.32	4.56	32.98	0.43	1.23	3.34	. C	9.00	₹.	47.09	2.09	0.05	0.67	0.36	58.08	0.09	33.36	0.12	3.10	1.87	0.22	0.07	0.28	0.22	1.81	2.18	90:0	20.42	11.62	6.38	4.28	0.14	0.25	1.79	
	AveBPH		0.59	0.81	5.93	0.08	0.23	990	3 ;	1.72	1.48	9.71	0.44	0.01	0.14	0.08	12.74	0.02	7.38	0.03	0.70	0.43	0.05	0.02	90.0	0.05	0.42	0.51	0.0	4.96	2.86	1.58	1.08	9.0	90.0	0.48	
Gen Bank	Accession	Number	R92362	AA235370	AA282273	AA148735	W16424	046449	01104	AA448484	N57002	AA425650	AA035745	N58138	R19878	T51539	AA425692	N57483	AA430545	H19446	N26556	W02426	N68139	AA128407	AA454570	N69499	N73563	H57585	T49325	AA485373	AA101878	N64798	AA487385	AA676604	N47012	AA883735	30000
1000			196168	687667	712950	504761	277662	25040	2000	782537	277596	773208	364865	247587	34888	72395	773204	277327	770086	61620	288343	296170	278504	526587	809526	292958	295997	205628	67397	841028	489837	284824	841507	882497	280155	1488788	3
			168	169	52	1	: {	4 5	2/2	174	175	176	#	178	178	68	<u>£</u>	182	£	2	<u> </u>	\$ \$	3 5	£ 5	2 2	3 3	ē	6	5	į	\$	\$	797	9	3 5	2	3

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor samples.

								A1767810181777	Fold/MPM122/A	Fold(MPM123/A F	old(MPM125/A	# of Tumor
	IMAGE	Gen Bank Accession	AveBPH	AveTumor	Fold(AveTumor/Av Fold(MPM120/A Fold(MPM121/A Fold(MPM121/A Fold(MPM120/A	old(MPM120/A veBPH)	Fold(MPM121/A	rolu(mr m 12-77)	veBPH)	VeBPH)	veBPH)	Samples
	Clone ID	Number						,	02.8	0.74	0.74	ო
1			č	200	3.89	3.91	7.78	48	0.0		9 40	en
202	723872	AA292583	5	9 1		6.41	6.16	96.0	0.35	જું :) t	
203	856174	AA630620	0.03	LT.0	80.0		7.01	2.57	0.65	0.88	A.0.	3 (
క్ష	782575	AA447522	0.36	1.39	3.86	6. 6	3.40	4.57	10.12	1.60	1.76	7)
205	428166	AA001870	2.28	8.77	3.85	F8.F		2.16	8.17	4 .00	3.75	က
208	586826	AA130857	1.17	4.49	3.83	2.76	5 6	9 0	7.80	2.29	4.30	က
207	730838	AA417012	0.02	90.0	3.81	7.50	84.0 66.0	, t	2.48	6.86	2.62	က
, e	837953	AA458578	7.85	29.78	3.79	3.76	5.93		6 6	5.74	6.57	က
2 6	503824	AA133959	0.12	4.0	3.78	1.62	0.09	0 7.7	189	0.76	1.61	က
3 5	208210	H65286	0.89	3.34	3.76	5.28	5.92	5.33 9.89	 	0.51	11.00	က
7 5	427998	AA001841	0.11	0.43	3.75	3.08	1.05	3.00	20.0	12.61	4.50	က
	423755	R01304	<u>48</u>	6.90	3.75	0.50	0.29	\$;	5 6	5.08	2.80	က
4 6	428483	A A 7 0 3 2 4 9	14.14	52.21	3.69	1.02	1.89	4 (0.63	3.32	4.58	က
2 2	443082	B71335	44.0	1.61	3.67	2.83	6.04	2.57	5.5	4.57	6.14	က
1 4 6	131599	R23727	0.16	0.60	3.65	5.05	2.39	2.70	6.1	6.33	5.22	က
3 6	240151	HR2708	2.33	8.49	3.64	2.83	0.64	28.L	- 000	3.62	2.52	က
2 5	483448	A A 707249	4	1.61	3.63	2.41	7.49	3.40	67.7	2.91	2.55	က
	204406	008477	1 05	3.81	3.61	3.93	4.00	2.14	• 6	1 20	2.78	က
917	671.07	730474 707448	720	88.0	3.55	6.64	4.18	2:20	4.29	2	2.81	က
ALZ	CI /071	611708	4 6	80	3.54	3.04	3.99	1.69	2.78		2.84	es
220	768179	AA424849	6.4	4 47	3.52	2.95	6.36	3.23	3.77	2.33		· e7
22	250678	H95976	4 .08	74:47	1 (9)	8	2.67	2.75	6.21	4.11	<u>è</u> :	
222	273489	N33263	0.09	0.32	3.52	9 9	3.80	3.48	7.92	1 .94	1.82	, C.
223	755785	AA496455	1.95	6.84	3.51	2.12	6.00	1.82	4.71	3.01	7.18	က
22	815047	AA465166	2.68	9.34	3.48	2.62	8. 5	3.77	7.25	2.34	2.03	က
225	384670	AA709333	4.17	14.53	3.48	3.11	2.41		365	2.26	3.29	က
228	782200	AA431964	1.53	5.30	3.47	2.48	6.32	5.7	6.25	3.48	2.49	က
227	52730	H29500	6.64	22.94	3,45	2.33	2.7) i	8.68	4.33	2.05	က
228	125311	R05810	1.92	6.62	3.44	1.07	2.84	6 6	÷	0.49	1.77	က
228	770880	AA434404	8.05	27.68	3.44	6.26	3.15	7.07	5 5 X	5.89	1.71	က
200	798288	AA460827	0.55	1.89	3.43	1.43	3.68	6.30	47.	1 12	1.39	က
3 5	204148	H55920	2.07	7.07	3.42	2.91	3.75	7.27	0 9	1 C	0.90	က
3 8	24.64	28000	5	90	3.41	9.82	3.38	1.53	3.80	9 4	0 X	m
737	5/5110	AA1/0055	5 6		3.40	1.86	1.13	5.78	5.41	51.1	9 6	~
233	325380	AA284285	5 6	3 6	3 30	5.38	1.97	4.32	1.12	1.58	9.0	· •
23	434822	AA703114	0.80	7/7		4 98	4.75	2.85	4.55	1.19	<u>.</u>	3
235	1049033	AA778675	1.87	6 .30	9,00) } }	7					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

					Seldimbation Foldimpation Foldimpation Foldimpation Foldimpation Foldimpation Foldimpation		A/10M104/A	Fold(MPM124/A	Fold(MPM122/A	Fold(MPM123/A F	old(MPM125/	s of Tumor
	IMAGE	Gen Bank Accession	AveBPH	AveTumor	Fold(AveTumor/A\	/ Fold(MPM120/A veBPH)	veBPH)	veBPH)	veBPH)	veBPH)	VeBPH)	seiders.
	Clone 10						90	2.99	5.61	3.78	1.72	
238	882483	AA676598	2.75	9.30	3.38	1.34	0 1	*	3.10	10.17	4.12	က
237	767130	AA424535	0.49	1.67	3.38	1.02	4.0	· · ·	7.41	0.36	2.21	က
8	126321	R06479	0.03	0.10	3.38	2.12	4 (. 44.6	2.89	8.21	0.15	က
239	230440	H75328	0.07	0.22	3.37	1.97	3.5/	, c	5.02	2.93	1.99	က
240	624271	AA181179	6.78	22.81	3.36	3.53	3.85	4.43	0.68	0.96	0.68	က
241	123265	R00284	2.30	7.72	3.35	4.22	9.13	1.43	1.48	3.53	5.19	က
242	357236	W83638	0.72	2.40	3.35	5.76	80.7	 88	0.96	3.48	2.80	က
243	123331	R00479	1.03	3.43	3.33	2.37	4.72	 	7.73	3.58	1.21	ო
2	240694	H78134	1.85	6.17	3.33	2.79	5.03 10.03 10.03	5.53	4 .	2.81	1.19	ന
245	347182	W80611	0.02	0.07	3.32	0.45	5.03	, e	4.13	2.87	0.31	ო
248	52092	H23232	0.03	0.11	3.31	1.94 4	ου 4 ου4	3 17	4.45	2.72	2.93	က
247	160628	H25019	0.41	1.36	3.30	2.02	80.4 84.6	4 6	1.55	3.28	3.34	က
248	1505294	AA905838	1.1	3.66	3.30	1.98	2.43	17:7	6.75	1.98	1.68	က
248	262053	H98812	7.69	25.26	3.28	2.38	3.42	0.40	2.34	2.25	5.58	က
250	741067	AA478436	17.59	57.78	3.28	2.99	3,38	r ac	4 85	1,83	3.72	ო
251	490178	AA121266	3.50	11.46	3.28	2.64	46.4	2.28	196	2.82	4.27	က
252	208741	H63175	0.09	0.30	3.26	2.00	19.4	3.64	, se	4.86	3.07	က
253	416611	W86468	3.63	11.85	3.28	2.01	2.14	2.13	3.05	3.65	6.62	က
254	416092	W85881	0.50	1.62	3.26	2.00	2.31	S 69 6	1.63	6.94	5.05	က
255	221561	H92525	0.55	1.79	3.26	1.18	1.35	3.80 3.80	2.95	5.14	7.79	က
258	223274	H86518	90.0	0.17	3.25	3.24	0.19	 	1.74	3.08	1.09	က
257	44387	H06525	1.17	3.78	3.24	6.67	2.69	<u> </u>	3.80	2.05	5.34	က
258	110787	190639	0.25	0.80	3.23	2.49	3.52	2. c		3.80	2.03	က
259	121948	T97762	4.17	13.48	3.23	1.59	4.12	7.45 	2.93	3.17	2.10	က
280	782812	AA448251	90.0	0.18	3.22	4.87	9.00	787	3.29	2.99	4.45	က
261	145513	R77718	0.67	2.16	3.22	1.90	10.4		3.83	2.74	3.25	က
262	435319	AA699931	6.14	19.70	3.21	2.30	5.27		5.80	5.20	0.95	က
283	210698	H66877	1.41	4.52	3.21	1.59	2.24	0.1	3.20	2.11	6.62	က
28	197067	R83401	0.36	1.16	3.19	1.88	1.56	5 6	2.00	3.26	2.91	က
265	416374	W86202	3.18	10.13	3.19	4.89	3.63	C+:3	88.0	1.08	3.06	က
288	1422794	AA827405	2.02	6.43	3.18	2.60	3.81	7.00	80.8	3.23	0.45	က
267	137017	R35665	0.03	0.08	3.18	1.47	12.4	2.53	3.68	3.77	2.82	က
268	267634	N34117	0.21	0.66	3.18	2.32	5.00	243	1.60	1.72	4.54	က
269	433253	AA699427	5.61	17.79	3.17	8. 8	ñ ŧ	e i				

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumbr sample or 3 fold or higher in at least 2 tumor samples.

1	Gen Bank			Eold/AvaTimor//	EsidiaveTimoriav Fold(MPM120/A Fold(MPM124/A Fold(MPM124/A Fold(MPM122/A Fold(MPM123/A Fold(MPM126/A # of Tumor	Fold(MPM121/A	Fold(MPM124/A	-old(MPM122/A	Fold(MPM123/A F	old(MPM125/A	# of Tumor
Accession		AveBPH	AveTumor	eBPH)	(HABPH)	veBPH)	veBPH)	veBPH)	Vestri		
Number				•	•	30.0	234	2.32	2.73	4.79	က
AA872692		7.39	23.26	3.15	3.45	3.20	. uc	136	3.88	2.18	ო
N26026		0.51	1.59	3.14	1.46	6.62	0.0	23	2.11	2.80	က
T61899	_	15.17	47.49	3.13	3.66	4.92	08.5	- c	3.41	4.13	က
AA427978	8	1.12	3.51	3.13	2.45	3.57	C 67.7	200	4.36	3.37	က
AA485676	. 82	2.17	6.72	3.09	1.06	2.45	0.70	1 2 3	5.53	2.34	က
N27935		99.0	7 .07	3.09	3.94	3.31	GL.2	- L	138	1.27	ო
AA702480	. 8	2.65	8.16	3.08	3.50	2.77	8. 6 4. 6	2 6	3.43	2.39	က
AA460557	2	0.48	1.46	3.07	2.06	2.12	3.31	6000	17.	4.45	ო
AA682831	<u>ج</u>	0.48	1,47	3.05	1.65	4.32	2.32	0.00	7	3.82	က
H71242		1.92	5.87	3.05	2.21	3.82	1.69	S .		3.78	က
A & 4.00598	, 8	0.49	1.50	3.04	4.39	4.19	1.69	CA:	27.4	19.1	က
TR3245	} <u> </u>	12.72	38.66	3.04	1.01	1.73	3.23	79.4	40 +	3.69	ო
A A R R R 739	, g	1.89	5.73	3.04	2.12	4.97	1.27	4.24	7.	3.08	က
2000N	2 2	1.72	5.20	3.03	2.45	3.22	5.41	2.32	07.0	4	က
R96561	. 19	0.51	45.	3.03	2.99	4.10	3.93	3.02	3 55	1.06	က
R58877	; =	19.83	60.07	3.03	2.41	3.80	2.96	0 70	4 +	1.32	က
44459879	679	1.63	4.94	3.03	3.06	5.11	2.63	P. 4	- c	2.42	က
W86779) a	0.57	1.71	3.02	1.77	4.00	2.37	70.4		1.97	က
H04262	: 2	1.16	3.50	3.00	1.70	2.11	3.21	6.15		2.18	ю
AA022498	8	0.03	0.08	3.00	2.52	0.38	4.22	3.89	3 6	5.43	ო
NOACA	3	90.0	0.17	3.00	1.60	3.79	1.68	27.7	89.0	2.56	ന
A A 0.7487	, [1.05	3.15	2.99	1.72	3.77	3.39	3.80	6.00	 0.80	. es
AARRADER	. 4	17.95	53.54	2.98	3.39	3.07	2.81	4.98	7.03 10.04	12.7	· «
003000	3 2	-	0.33	2.98	1.54	3.21	1.65	5.62	5.		۰ ۳
A 4 6 6 6 7 0	2 6		2 79	2.98	4.82	3.25	2.81	3.28	1.52	7 7 7) e
			3.41	2.98	5.58	4.11	2.12	,	3.10		. «
CCC70/W		2 3	, F.	800	2.82	3.27	4.53	1.90	1.93	3 ·	.
AA877997	.68	5 .5	0	90.6	135	3.39	3.86	2.74	3.89	2.47	n (
H51039	39	0.22	0.65	CR.7	2 1	338	1.71	2.34	0.80	3.84	r>
AA863449	¥	1.15	3.38 8.	2.94	0.07	900	2.31	3.06	2.04	3.43	က
H48278	28	0.83	2.42	2.93	2.70	1 6	3.44	1.10	1.06	0.29	က
H99694	2	0.75	2.19	2.93	3.82	00.7	. · ·	23.4	77.7	2.22	က
AA001978	976	0.05	0.14	2.91	0.99	3.14	9. Q	706	3.55	1.25	ო
N95187	87	0.35	1.01	2.91	2.20	4.92	3.17	7.07	3 7	141	က
AA458959	929	90.0	0.17	2.90	1.34	3.15	1.41	o o	ř	<u>.</u>	
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Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold of higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

	IMAGE	Gen Bank Accession	AveBPH	AveTumor	Fold(AveTumor/Av	Fold(MPM120/A veBPH)	Fold(MPM121/A veBPH)	Fold(MPM124/A vebph)	Fold(MPM122/A	veTumor/Av Fold(MPM120/A Fold(MPM121/A Fold(MPM122/A Fold(MPM123/A Fold(MPM126/A # of Tumor RPH) veBPH) veBPH) veBPH) veBPH) veBPH) veBPH)	Fold(MPM125/A veBPH)	# of Tumor Samples
	Cione 10	Number						3.29	1.57	3.43	4.63	- ო
Š	1460820	AA889785	3.62	10.49	2.80	2.35	7. 7.	3.89	2.10	2.31	2.77	ო
305	26259	R20547	0.87	2.50	2.89	3.08	4.0	8 5	1.23	1.00	6.23	ю
308	366154	AA062813	0.0	0.03	2.85	3.87	3.77		1.52	0.39	3.94	က
307	282104	N51498	0.93	2.66	2.84	2.66	0. i	0 C	1.78	1.21	1.74	က
308	344308	W70147	0.46	1.31	2.81	3.18	74.6	9.30 4.80	3.52	2.33	2.40	က
308	45525	H08725	2.19	6.17	2.81	2.20	3.24	 	2.46	3.30	0.95	က
310	196303	R92435	9.0	0.11	2.80	2.56	5.0.4 5.0.4	, c	3.41	1.52	2.19	ო
311	267495	N25262	1.59	4.40	2.77	3.68	3.24	3.45	1.24	1.90	1.53	က
312	757198	AA496133	0.65	1.78	2.75	3.73	, v. v.	3.40	3.07	2.93	0.52	က
313	204122	H61886	1.31	3.59	2.75	2.36	4.23	, t	1.18	3.64	3.82	က
314	839545	AA489791	0.42	1.1	2.73	5.37	68.0		2.89	4.45	4.47	က
315	809694	AA454702	0.02	0.05	2.70	0.54	0.56	5.50 2.40	1.13	3.56	3.32	ო
316	126234	R06362	0.30	0.81	2.69	2.13	3.70		3.26	3.22	2.72	ო
317	280407	N51563	0.97	2.62	2.69	1.61	04.6 04.0	 	3.46	3.11	4.17	က
318	758284	AA404231	0.78	2.08	2.67	1.82	2.31	 	2.04	1.07	2.09	ဗ
319	150314	H00817	2.18	5.81	2.67	3.69	3.80	3.52 00.8	. 66.0	2.89	3.97	က
320	288936	N62696	9.0	0.11	2.67	1.59	6.47	1 79	4.52	0.21	5.08	ဗ
321	757440	AA437226	0.05	0.13	2.67	0.21	91.4 90.0	2 - c	3.33	3.56	1.30	က
322	176554	H45289	0.87	2.28	2.64	1.89	2.02		1.85	3.26	3.01	က
323	298057	N69989	0.68	1.79	2.62	2.03	3.01		66.6	0.70	3.96	ო
324	755299	AA496359	1.69	4.39	2.60	4.47	0.52	t.	2.30	1.62	1.73	က
325	725321	AA291749	0.87	2.27	2.59	3.21	3.07	5.01	0.72	3.58	4.52	က
326	1467831	AA883431	1.17	3.02	2.59	1.16	1.22	7:35	986	3.01	3.31	က
327	120097	T95151	0.60	£.	2.58	5.09	L0:2	† † ¢	790	3.72	4.51	က
328	159455	H15746	0.02	9.0	2.58	0.62	3.70		232	3.41	3.22	ო
328	1460696	AA868802	6.57	16.82	2.56	1.39	3.08		7.0	0.24	4.83	က
330	757368	AA437126	0.02	90.0	2.55	3.92	2.20		300	4.79	4.4	ო
331	450612	AA682452	0.74	1.89	2.55	1.17	0.96	0.92	1 17	4.50	0:30	က
332	342543	W88141	0.05	0.0	2.53	4.17	44.	0 6	137	1,19	1.85	က
333	503741	AA131466	1.05	2.65	2.53	3.31	3.85	3.58	. T	3.28	1.79	က
33	784130	AA432075	17.73	44.77	2.52	3.17	4.60	0.76	<u>.</u> .	0.31	3.14	က
335	325370	W52208	0.03	0.08	2.51	3.68	3.87	78.7	9 6	1.37	3.87	က
338	288663	N62394	0.78	1.95	2.50	3.07	3.96	C. 63	90 E	7.02	90.0	က
337	344359	W73377	0.07	0.18	2.46	0.90	3.52	<u>t</u>	3			
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Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor samples.

	-	•				THE POINT PRINT PR	V/POPILIZATION	Poly(MPM124/A	Fold(MPM122/A	Fold(MPM123/A F	old(MPM125/A	# of Tumor
	IMAGE	Gen Bank Accession	AveBPH	AveTumor	Fold(AveTumor/A	v Fold(MPM120/A veBPH)	roid(MFM1211A veBPH)	VeBPH)	veBPH)	veBPH)	veBPH)	Samples
	Clone ID	Number			(u.a.			70	440	1.23	1.35	က
000	10004	D48244	1.15	2.79	2.41	3.60	3.57	\$	2 4	50	3,55	က
3	- 6000	1450000	7	0.35	2.40	3.16	3.19	1.82	60:	, K	2.32	က
200	428825	AAUUSOUS	<u> </u>		077	3,61	3.23	0.26	4.12	6.0	7	্ ক
8	258242	N30655	0.62	P.		08.0	4.21	1.0	1.52	3.40	<u> </u>	•
34	255897	N27366	0.09	0.20	2.31	8 6	3.24	1.70	3.19	3.01	1.39	, co
342	452068	AA707125	0.88	2.03	2.30	87.T	, c	0.16	3.57	0.38	3.18	က
343	565779	AA135870	20.69	48.64	2.25	2.43	0.0	2.0	3.30	0.24	4.05	က
7	128506	R10154	9.0	0.09	2.15	1.20	3.8/	7 0	2	0.88	0.82	က
5	839048	AA487505	0.36	0.76	2.12	4.12	3.45	0.42	1.78	4.85	73.83	~
38	280763	N50563	0.03	0.50	14.36	0.58	2.77	2.30 80 80	9.24	42.46	32.88	8
3	590853	AA158162	0.0	0.18	13.25	0.74	0.74	- e	136	2.46	58.40	7
	951101	AA620472	0.02	0.22	11.30	1.40	0.82	5.57 8.00 8.00	20 07	1.07	3.31	8
3	377610	AA055833	0.13	<u>ب</u> بج	10.02	1.46	2.83	2.30 2.30	2.67	34.78	5.82	7
350	50787	H16833	90.0	0.50	7.90	1.60	0.16	6.30 234		1.1	2,44	7
35	137984	R63085	0.29	2.25	7.79	8.43	33.82	6.5	943	32.14	1.71	~
352	856585	AA669222	0.12	0.94	7.77	1.46	0.70	D	0.24	35.70	0.66	7
, E	358990	W92263	0.06	0.49	7.64	1.97	2.74	6. 4. 0. 4.	3.78	1.47	2.84	7
2 2	430102	AA010328	0.10	0.75	7.47	33.12	2.43	A	9	970	2.55	8
	454477	100837	1.19	8.16	6.87	7.32	29.35	1.05	B 7	5 E	1.40	8
000	774101	T02031	74.0	3.19	6.81	15.33	19.31	2.50	\$ 5	5	0.36	8
8 1	55.0	11001	20.0	1.88	6.69	13.67	22.88	0.57	1.63		5.36	84
3	02020	766974	800	0.53	6.43	1.07	0.12	0.12	0.52	01:00	18.17	~
3	770808	AA0004U3	2000	1 39	8.27	0.88	0.41	0.38	1,35	9.49	20.50	. ~
200	000674	AAUUSTBO	1 8	33	5.48	6.98	2.35	1.09	0.55	06.1	72 03	
8	264449	NZ1233	9 6	30.7 80.7	5.45	2.04	2.77	1.77	2.77	11.30	5.03	
	754358	AA436142	0.60	3 5	5.18	1.16	22.31	1.49	3.68	L8:0	9 0	۰ ،
362	07790/	A424800	3 6		5.17	4.05	23.41	98.0	1.45	5. J		, ,
383	303139	N90806	0.02	- 7:7	 	0.47	1.51	0.94	20.25	2.06	05.0	
8	757165	AA443950	30.0	2 8	20:0 4 88	7.86	14.43	2.52	1.47	0.61	2.40	
365	565235	AA136125	18.25	99.00	9 6	3.70	1.71	<u>4</u> .	0.22	2.88	18.58	v
366	247103	N53959	0.05	0.22	79.4	 A	2.58	4.82	2.82	14.34	2.86	7
387	234376	N28268	7. 7.	14.16	IO.4	? c	1.45	2.24	3.15	2.40	18.90	74
368	758405	AA482119	9.0	0.21	4.73	0.22	2 5	0.55	0.22	9.85	2.80	8
369	47355	H10981	0.10	0.48	4.65	14.30	2 7	11.68	1.29	2.30	2.82	~
370	377898	AA777098	3.48	15.28	4.38	5.72	£.2.5	123	1.41	1.47	0.93	8
371	745072	AA626275	0.45	1.95	4.32	6.41	•	2				
							7					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

							of Tumor Fold(MPM123/A Fold(MPM123/A Fold(MPM125/A # of Tumor	2 4/7072	AldinDM1221A	old(MPM123/A	Fold(MPM125/A	, # of Tumor
	IMAGE	Gen Bank Accession	AveBPH	AveTumor	Fold(AveTumor/Av F	old(MPM120/A	Fold(MPM121/A	rold(mrm14477 veBPH)	veBPH)	VeBPH)	VeBPH)	Samples
	Clone ID	Number						4 07	1.48	0.06	14.64	-
372	79763	T63981	0.16	0.71	4.32	1.17	84.7	5 6	1 79	0.93	4.17	7
373	429424	AA007687	0.03	0.14	4.20	2.13	15.88	0.50	20.7	0.56	0.11	8
374	265655	N22836	0.09	0.39	4.17	16.96	1.24). (6. 0	13.79	4	1.42	8
375	1606315	AA991196	0.72	3.01	4.16	2.17	2.93	3.40	0.52	0.21	1.46	8
376	1389018	AA855158	0.05	0.20	4.11	11.10	10.86		2.67	1.88	1.99	8
377	283335	N51441	0.18	0.72	4.00	1.22	10.53	9 6	750	0.92	10.57	7
378	284931	N71861	0.02	0.07	3.99	2.68	8.36		2	0.36	15.08	7
379	1031203	AA609982	0.03	0.11	3.96	1.17	0.36		4.17	0.30	16.49	7
380	758360	AA404290	0.03	0.13	3.95	0.30	0:30	2.50	2.97	1.40	10.58	8
88	178569	H49053	0.05	0.20	3.95	3.16	3.00	2.58 1.08	1.67	2.16	12.53	8
382	154477	R54855	1.01	3.95	3.92	2.32	6/:1	9 6	2.22	1.03	0.56	7
383	1048998	AA778636	0.14	0.57	3.91	10.73	5.8c	7.80 00 00	0.93	3.45	1.09	8
38	129589	R16541	4.0	1.66	3.79	2.77	13.43	60:- C	27.2	1.71	9.27	7
385	1055636	AA620927	0.19	0.72	3.75	3.32	2.92	6.08 80.00	8 45	90.9	1.64	7
386	461699	AA682274	10.27	38.51	3.75	0.81	2.56	9. 80 8. 80 8. 80		5,65	6.87	8
387	50692	H17626	0.31	1.18	3.74	1.87	2.97	9 6	11.37	0.71	0.71	7
388	362926	AA018980	1.07	4 .00	3.74	2.76	5.27	20.0	ā	0.20	10.81	8
389	122178	T98615	0.03	0.11	3.69	0.65	8.21	t 6	180	0.52	0.85	84
390	246808	N53214	0.56	2.06	3.69	6.45	12.29	0.2.1	88	5.39	1.99	7
391	191866	H40350	0.61	2.24	3.68	2.33	6.76	20.3 84.0	0 82	0.72	1.33	8
392	155072	R71393	0.30	1.07	3.62	7.00	9.56	5 5	2.82	7.51	3.33	8
383	190325	H29897	0.70	2.51	3.59	2.78	2.7		7.62	1.52	6.75	8
ğ	502438	AA134862	1.10	3.8	3.58	1.28	2.72		1.03	10.16	1.08	7
395	293689	N69675	0.01	0.03	3.57	6.95	1.00	2.0	10.40	1.10	2.70	7
396	1486752	AA912071	0.70	2.48	3.57	1.94	9.07	7.88	1.98	1.85	2.87	81
397	510136	AA053239	0.03	0.10	3.52	6.17	4.48	27.5	7.18	5.86	2.39	п
398	878284	AA670280	2.85	9.83	3.45	1.42	96		2.62	2.54	1.99	N
388	321470	W32303	1.07	3.68	3.45	3.79	67.7	£ ;	1.12	7.17	1.71	7
\$	287866	N62332	1.24	4.26	3. 4.	2.88	at c	27.0	7.36	2.52	1.53	71
5	258300	N30680	5.10	17.12	3.36	2.43	 	3.67	2.29	2.53	6.91	8
405	1048646	AA608893	0.92	3.08	3.35	1.96	7 7 7	, 1	6.82	1.99	1.86	8
4	289025	N59826	0.55	1.82	3.35	2.20	87.6 7	- e	2.13	1.50	1.49	8
ş	291706	N73480	0.53	1.77	3.33	4.14	1.81	4 6	2.61	1.62	4.45	8
504	1031568	AA609304	0.08	0.25	3.30	6.38	/R.7	<u>:</u>	i			
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The state 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

		75.00	•		September 1 Pold MPM124/A Fold MPM124/A Fold (MPM122/A Fold (MPM123/A Fold (MPM125/A # of Tumor	A)004HBBH)FI	Eold/MPM121/A	Fold(MPM124/A	Fold(MPM122/A	Fold(MPM123/A	-old(MPM125/A	# of Tumor
	IMAGE	_	AveBPH	AveTumor	Fold(Ave Tumor/Av	VeBPH)	veBPH)	veBPH)	veBPH)	veBPH)	VeBPH)	Calming
	Clone	Number					ğ	2.77	2.09	2.39	1.81	8
904	383676	AA679116	0.13	0.41	3.28	7.55	9 6		4.55	5.37	2.84	7
404	310493	W31074	6.22	20.37	3.28	2.57	2.10	700	2.76	0.15	0.69	8
408	84713	T74257	0.07	0.21	3.27	6.85	0.10	10.73	12.99	4.14	0.35	8
604	219955	H81817	0.19	0.61	3.24	0.75	0.51	0.73	2.27	4,35	6.93	7
410	295741	W02483	0.35	1.13	3.23	2.04	2.14	6 6	8.52	1,58	5.09	8
±	796646	AA460115	9.12	29.44	3.23	1.38	0.93	9 C	3.86	1.48	10.68	8
412	44253	H05818	90.0	0.21	3.23	1.53	1.2.r	7.62	2.09	3.85	2.98	8
4	139051	R62926	0.44	1.43	3.23	2.91	2.95	, c	10.50	3.02	1.53	8
4	454543	AA677023	0.83	2.67	3.22	0.89	1.39	2.00	5.55	7.72	2.25	8
415	48315	H15368	0.22	0.71	3.21	0.69	1.60	- C	1.50	3.24	0.62	84
418	346612	W74462	1.22	3.89	3.17	2.96	98.6	. a	55.	1.43	1.53	8
417	263836	H99766	1.11	3.52	3.17	2.49	4.47	, t	443	2.52	96.0	7
418	376793	AA046883	0.72	2.28	3.15	7.40	8c.r	- 4 B	7.85	4.94	2.01	8
4	283617	N52875	1.18	3.69	3.13	0.71	1.64		53	0.65	1.06	7
420	1684274	A1003028	1.45	4.54	3.13	5.71	7.10	6.73	- c	6.73	1.25	8
4 24	784212	AA446864	1.91	5.92	3.11	2.46	2.18	2:50	, c	0.17	9.50	8
733	41214	R56898	0.03	0.11	3.11	0.98	3.88	80.	6.35	0.37	2.46	8
3	384688	W96197	0.03	0.08	3.10	7.63	1.12	4.50	2.04	330	2.00	
2 2	149013	R82289	7.60	23.51	3.09	0.77	1.07	3.65	8.67	2.3	4.03	8
202	392399	AA707835	0.52	1.59	3.09	2.36	2.94	2.93	gr.2	7.0	0.99	7
100	347615	W81526	0.07	0.20	3.08	12.89	0:30	0.86	8.7.0 1	; 4	8.48	7
127	1030779	AAB09004	0.52	1.61	3.08	1.29	1.48	1.52	9. %	9 8	1.60	8
2	364555	AA022801	18.	5.67	3.07	5.36	2.73	1.56	1.33	9 68	1.28	7
4 28	743688	AA629324	1.27	3.90	3.07	2.60	2.21	5.26	5.1.6 4.0f	6 4	0.24	8
63	141726	R69584	90	0.13	3.07	8.46	2.87	3.48		7	4.82	8
5	786147	AA480986	2.98	9.08	3.07	2.49	2.65	2.36	1.7	100	5.35	8
432	813501	AA456078	0.60	1.85	3.07	1.76	2.60	4.83	27.1	418	4.86	8
533	138047	R53757	0.35	1.07	3.04	2.90	1.81	2.18	5.55 5.50 5.50 5.50	7.13	1.15	7
\$	22800	T87235	0.12	0.37	3.04	3.96	5.88	1.86	2.82	0 7. C	4.	8
435	201818	H48318	0.16	0.48	3.04	2.44	5.47	4.57	1.88	2.14	2.50	~
85	345781	W72871	0.85	2.80	3.04	2.94	5.24	1.48	2.88	- e	107	~ ~
12	42747	R61847	0.22	0.67	3.03	1.17	1.54	1.10	08.8	5.50	4.58	~ ~
£	743230	AA400234	8.50	25.66	3.02	2.49	2.16	1.72	2.64	 	£73	. ~
65	261580	H98694	8.42	25.35	3.01	2.74	2.82	1.36	2:36	e F		
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Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

		Gen Bank					COLUMBIA 1941A	Fold/MPM124/A	Fold(MPM122/A	Fold(MPM123/A F	old(MPM125/A	# of Tumor
	IMAGE Clone ID	Accession	AveBPH	AveBPH AveTumor	Fold(AVe I umor/A eBPH)	V FOIG(INFINIZOR V	veBPH)	veBPH)	veBPH)	veBPH)	veBPH)	Samples
;	00000	Jeguina Jeguina	7	72.0	3.04	3.08	1.37	8.10	2.01	2.27	1.23	7
3	979/05	1214W	. c	5 7		1.74	5.33	1.83	2.67	2.63	3.85	7
‡ :	89/10/	AA282655	77.	2.00		2.24	4.24	4.46	1.57	2.77	2.74	8
7 :	/04CR1	K82103	3	1 1	986	1.42	2.73	1.25	1.76	5.99	4.80	7
? :	400.004	AA6/6820		70.7		4.17	6.87	1.66	1.67	2.47	1.08	8
:	100238	A4075804	. e	£ ¢	2.87	2.12	6.90	1.16	4.61	0.14	2.91	~
F 3	20000	1470834	7 2	22.53	2.87	2.25	2.76	1.65	2.96	4.30	3.91	~
\$;	800045	W/8834	6.0	20:22 8 24	2 07	2.52	2.84	2.73	4.48	3.12	2.14	74
Ì	143/90	K/6/82	80.7	7. 6	90 6	1 02	2.38	1.71	3.88	6.14	2.52	7
1 3	50530	H17919	C.3	5 9	9. so	30.0	6.36	2.48	4.67	0.81	1.18	7
2	208082	H62563	77.	12.40	2.90	0 1 0	1.59	10.45	1.71	0.15	3.62	7
4 8	480525	AA126468	0.07	0.20	CR.7	2 6) (a	204	2.25	0.12	2.05	7
4 5	264157	N20602	0.09	0.26	2.85	4.52	0.60	10.7	06 4	3.21	2.23	8
452	133440	R27450	1.72	5.05	2.94	2.28	2.49	0. i		, t	2.83	2
453	416889	W86832	0.52	1.52	2.82	1.91	5.15	4.77	1.53	6) : :	
25	271076	N29918	2.08	6.09	2.92	4.05	4.90	2.55	2.04	7.07	6.30	4 (
455	452320	AA704777	1.07	3.13	2.82	2.22	2.90	2.16	3.07	2.67	o (۰ ،
458	814798	AA455235	25.12	73.30	2.82	3.83	7.12	1.66	2.07	1.26	S	7 (
457	417908	W80520	5.97	17.42	2.92	2.62	6.06	2.91	4.81	0.62	0.47	79
, K	745283	A A R 2 K R 7	5.50	16.00	2.91	2.17	1.35	1.50	2.45	5.86	4.13	2
3 2	K0772	146803	10.55	30.4	2.89	3.09	5.75	2.12	2.83	1.37	2.15	7
	7 000	710003	3 6	90 +	88 6	2.55	3.73	3.99	2.18	2.15	2.70	7
ş ;	90916	H18936	8 6	08 c		1.27	2.47	2.59	4.88	4.06	2.03	8
;	400000 101000	133	5 4	8 6	2.87	5 08	3.10	4.25	2.58	2.78	2.46	8
70,	20016/	709449	- d	0.43	2.87	2.66	3.84	3.59	2.24	2.73	2.16	8
3	124081	AT LOOK	0.55	5. 4.	2.87	2.46	3.04	2.60	5.16	2.82	1.11	7
¥ 4	45597	107034	1 69	4.82	2.85	2.11	5.17	3.34	2.83	1.61	2.04	~
8 4	721284	000000	5	0.13	2.84	2.74	1.81	3.99	1.81	2.21	4.51	~
3 4	48055	HORRED	97.0	2.08	2.84	0.41	0.65	0.73	4.40	10.07	0.75	7
. 8	ASASA	AA180113	29.81	25	2.83	2.15	2.80	3.18	6.89	1.20	0.76	~
3	276450	N34824	0.47	1.32	2.82	6.04	3.19	1.65	1.41	2.48	2.16	~
67.	480435	AA677561	3.97	11.17	2.82	2.79	3.27	2.11	2.34	4.55	1.85	7
	41825	R54109	0.07	0.21	2.82	0.83	4.13	2.29	2.36	0.14	7.15	7
	682058	AA258458	1.86	5.21	2.80	4.23	4.71	2.40	2.75	0.67	2.03	64
£	754408	AA436187	0.03	0.09	2.78	5.62	0.31	1.97	2.16	4.95	1.67	71
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Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold of higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

	IMAGE			1	Fold(AveTumor/Av Fold(MPM120/A Fold(MPM121/A Fold(MPM122/A Fold(MPM123/A Fold(MPM126/A # of Tumor	Fold(MPM120/A	Fold(MPM121/A	Fold(MPM124/A	Fold(MPM122/A	Fold(MPM123/A	Fold(MPM125/A	# of Tumor
	Clone ID	Accession	Аувври	Ave i umor	•BPH)	VeBPH)	veBPH)	VeBPH)	(Bage)			
į		reguna	6	20.24	9.78	2.12	2.47	2.22	3.18	4.00	2.68	7
4/4	4584	H28215	13.78	10.00	2 }	1 48	3.14	4.59	2.82	1.71	2.89	8
475	291222	N72196	90.08	0.23	77.7	 	2.91	2.83	0.95	2.38	4.21	ч
476	769857	AA430367	1.25	3.40 0.40	2.70	9 0	2.80	4.86	2.49	2.42	3.18	7
E#	195875	R92216	0.59	\$ 6	2.70	, c	2.17	0.86	4.74	2.36	5.84	7
478	731048	AA421275	0.02	6.6	2.76 9.78	4.00 36	1.66	2.64	3.65	1.52	2.70	8
479	812965	AA464600	0.87	2.41	. 07.7	5 4	, y	2.60	3.44	1.72	1.19	8
480	809789	AA454758	0.17	0.46	2.76	1.60	3.74	2.54	0.51	2.26	6.97	8
2	289513	N62745	0.02	0.05	2.75		173	3.12	1.74	1.08	6.80	7
482	327239	AA284307	0.50	1.37	2.73	4.00 2.20	. 4	3.71	2.22	1.48	2.78	7
\$	23588	R38369	1.07	2.98	2.75	27.7	3.42	2.33	2.85	1.79	4.41	7
4	153617	R48833	1.27	3.49	2.73	7.75	4.33	2.24	4.29	1.74	2.98	8
485	491435	AA150435	0.28	0.78	2.75	0.83		88 0	0.55	8.83	4.	7
486	236282	H61193	0.12	9,34	2.75	1.13	0.00 0.00	26.5	3.17	0.83	7.34	8
487	781342	AA448390	0.08	0.22	2.75	2.79	0.80	00.	. 60	1.28	1.51	8
488	786525	AA452113	0.81	2.21	2.74	7.49	3.35	0.82	c	20.4	1.83	и
489	742082	AA405769	0.13	0.35	2.74	2.57	4.19	6.30	2.3 80 80	4 23	2.31	8
8	39821	R53428	12.95	35.29	2.73	2.05	2.54	2.17	9 6	6 6	77.0	8
481	40338	R53072	2.38	6.50	2.73	2.54	4.72	4.82	2.58	50.5		۰ ۸
482	194638	R84398	10.30	28,05	2.72	2.21	1.67	49:	1.82	00.4	\$ F	
107	149406	H01516	3.58	9.74	2.72	1.80	2.57	3.46	4.97	1.74	//:1	,
7	74223	T48412	6.45	17.51	2.71	3.86	6.49	1.22	0.80	1.93	88.L	4 (
100	464708	A & & 77730 K		4	2.71	2.51	1.34	3.04	1.20	0.70	7.49	N (
6 4	428338	AA005153	130	3.51	2.70	1.67	3.06	2.89	2.48	3.61	2.51	7 (
} {	140000	A 4 6 6 2 F F 6	3 2	9	2.70	1.84	2.59	3.68	1.98	2.26	3.86	8
	281204	H08218	0.05	4.0	2.70	4.16	4.07	1.82	2.64	1.18	2.32	N (
9	1838071	A1047332	33	3.57	2.69	1.60	2.33	3.00	5.58	4.99	1.6 6	N (
	202200	NERSO	0	5.39	2.69	3.78	4.41	1.27	2.53	1.43	2.71	2
3 3	000707	A A B 70288	8 6	10.81	2.69	2.74	3.62	1.27	1.97	4.20	2.33	N
5 6	107070	201020	, v	40 41	2.68	1.36	3.04	2.45	1.92	4.73	2.60	~
8	184304	1,000H	2 e		2.68	0.63	2.98	7.53	3.18	0.92	0.82	8
3 3	810328	AA484443	0.28	0.75	2.67	3.26	2.72	2.12	2.74	2.16	რ გ	%
	80414	U48743	20.0	5.53	2.67	2.59	6.02	0.45	5.94	0.35	0.70	N
3 5	743734	A A B 20257	28	0.76	2.67	2.85	1.51	4.67	0.86	4.27	1.88	~
202	452047	AA707118	14.0	1.08	2.67	1,41	6.37	1.25	3.28	1.41	2.28	~
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Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

	IMAGE	Gen Bank	AveBPH	AveTumor	Fold(AveTumor/Av Fold(MPM120/A Fold(MPM124/A Fold(MPM122/A Fold(MPM123/A Fold(MPM126/A # of Tumor - veBPH) samples veBPH) veBPH) veBPH)	Fold(MPM120/A	Fold(MPM121/A	Fold(MPM124/A	Fold(MPM122/A	Fold(MPM123/A F veBPH)	old(MPM125/A veBPH)	# of Tumor Samples
	Clone ID	Number			6 878)			7,7	1.76	3.89	4.18	2
508	361587	AA016980	0.88	2.34	2.85	2.25	2.02		. r.	3.31	2.24	81
509	283142	N51335	4.33	11.49	2.65	1.56	2.50	0.0		1.43	1.03	2
510	342721	W68291	0.58	1.53	2.65	1.63	2.24	5. C	2.40	0.20	2.38	8
511	530185	AA111969	0.31	0.83	2.65	4.40	3.73	80.2 00.0	. .	48 :0	0.68	2
512	395609	AA757588	0.03	0.07	2.65	7.24	3.66	2.08	2.49	4	3.00	8
513	898318	AA598828	2.48	6.52	2.65	3.03	4.78.		1.86	1.88	2.52	8
514	241787	H91721	0.16	0.42	2.65	1.72	3.98	- 6- - 6-	<u> </u>	3.28	1.73	7
515	1636108	A1015679	2.30	6.08	2.65	2.96	2.48	5.70 5.70	06.	2.76	3.37	7
516	201172	R98487	0.67	1.76	2.65	1.56	4.16	7.12	157	4.53	4.38	8
517	228290	H83865	0.70	1.86	2.64	1.33	2.40	t 80	1.19	0.26	8.46	7
518	342164	W63776	9. 2	0.10	2.64	0.60	0.20	9 6	2.86	3.04	0.98	8
519	49394	H15655	0.51	1.35	2.64	2.46	14.4	50.7 70.00	2 80	3.08	2.30	7
520	85320	171889	6.68	17.60	2.64	1.58	1.93	- 4 - 4	1.65	76:0	1.14	7
521	115143	T86708	5.38	14.12	2.63	3.65	6.57	5 6	88.	5.62	4.26	7
522	31969	R41994	0.74	1.96	2.63	1.11	50. 1.08) } }	1.03	1.50	8
523	362402	AA018449	9.88	25.98	2.63	2.80	4.67	7.77	4 6.	0.85	0.77	2
524	263049	N20046	0.35	0.91	2.63	1.09	4.25	8	2 2	2.21	2.76	7
525	31299	R42946	0.73	1.92	2.63	1.79	3.06	14:7	2 4	2.47	2.40	81
526	62112	T41077	3.92	10.27	2.62	3.22	2.98	94.	5.5	2.17	3.98	7
527	49728	H29198	0.70	1.83	2.61	2.66	2.18	6. to	 	2.40	3.25	7
528	295723	N66925	2.82	7.37	2.61	2.25	2.74	3.72	88	2.16	1.28	8
529	447715	AA702781	14.05	36.63	2.61	2.78	3.25	<u> </u>	2.53	3,64	3.38	8
88	898070	AA598779	0.92	2.40	2.61	1.58	2.75) ; ;	50.0	45.6	3.66	7
53	60540	T40531	1.12	2.91	2.60	2.02	2.15	£.4 4.5	08.6	2.10	2.06	8
532	288999	N62726	3.95	10.29	2.60	2.27	3.83	£ .	4.60	1.96	2.40	8
533	290346	N62293	1.53	3.89	2.60	1.93	3.19	<u> </u>	. E.	3.45	2.14	8
55	51865	H23187	0.05	0.12	2.59	2.52	1.93	07:	88.8	4.1	2.42	~
535	815124	AA481491	1.93	5.02	2.59	1.28	1.68	9 50 9 60 9 60	-	3.23	3.47	ч
538	428296	AA004944	0.60	1.58	2.59	1.85	2.54	7.00 00 6	0.82	2.24	0.98	7
537	51607	H19427	0.15	0.40	2.58	2.50	9.88 10.00		0.37	7.68	3.22	8
538	283461	N50647	90.0	0.14	2.58	1.74	2.31	9 6		0.14	0.92	7
539	283190	N51357	0.07	0.19	2.58	1.89	5.87	4.08 20.0	4 23	1.30	1.49	7
3	74511	T59024	10.71	27.60	2.58	3.63	2.79	2.02	0.68	10.78	0.24	7
24	300051	N78927	1.59	4.10	2.57	3.34	0.28	1	3			
							16					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold of higher in at least one fund sample or 3 fold or higher in at least 2 tumor samples.

	!	Gen Bank			Entrance Englimental Foldimental Foldimental Foldimental Foldimental Foldimental Foldimental Foldimental	Ald/MPM120/A	Fold(MPM121/A	Fold(MPM124/A	Fold(MPM122/A	Fold(MPM123/A	Fold(MPM125/A	# of Tumor
	Clore	Accession	AveBPH	AveTumor	Polician (ARB)	veBPH)	veBPH)	veBPH)	veBPH)	VeBPH)	Version)	e de la composition della comp
		Number		;	. !	ò	1.51	1.84	3.37	5.78	1.95	и
242	482354	AA705525	1.39	3.59	2.57	0.80	5 6	1.52	4.66	4.97	1.15	7
£.	243159	H94471	1.79	4.60	2.57	1.22	. 6. 6	2.87	3.13	1.36	1.13	8
\$	824896	AA488893	0.07	0.19	2.57	A	f F	3 10	1.45	5.51	2.23	7
545	884283	AA669750	9.33	23.95	2.57	1.33	0.70		2.31	3.57	1.65	8
248	260707	H97186	0.37	96.0	2.56	1.7	3.55	9 6	25	3.25	1.49	8
52	140197	R66101	0.29	9.74	2.56	2.80	B	6.08 6.09	333	17	1.17	и
248	344618	W74725	0.07	0.18	2.58	4.68	2.29	8.7.5 0.00	4 05	4.73	2.53	8
25	270535	N33228	0.49	1.25	2.55	1.19	2 8.	3.08	6. 1	25. 25.	3.87	8
550	811023	AA485380	0.22	0.56	2.55	1.72	1.10	97.1	<u> </u>	2.84	4.51	8
551	208413	H62162	0.34	98.0	2.55	3.23	2.89	0.80	5 6) i	2.72	8
552	290227	N62271	3.70	9.39	2.54	2.58	3.35		£.78	35.6	3.60	8
553	305677	N89973	1.10	2.79	2.54	0.77	2.86	86.	7.40	148	1.59	8
555	298554	N70592	2.03	5.15	2.54	2.50	3.69	1.78	7.40	 88	2.38	8
555	79000	T61938	6.02	15.26	2.54	3.54	2.61	3.14	00. 84	20:- A	1.66	~
558	321455	W45014	0.57	4.	2.54	. 4	4.26	1.05	7.20		3.89	7
557	1456625	AA864704	3.91	9.90	2.53	1.95	1.48	2.26	3.11	2.3E	0.95	- 8
558	796723	AA443140	0.14	9.3	2.53	2.23	3.90	1.93	† 6	, ;	2.36	8
559	85670	T62072	0.80	2.01	2.53	3.12	3.59	1.80	2.56	2	3	. 64
280	365738	AA025538	1.74	4.38	2.52	2.94	3.50	1.27	3.56	06:1		
5	489722	AA099593	3.24	8.15	. 2.51	5.69	1.78	3.26	1.83	3.20	2.32 2.05	۱ ،
582	1488074	AA889397	0.50	1.26	2.51	1.77	1.90	2.87	3.11	5.50 2.60		۱ ۵
883	252683	H88329	1.39	3.50	2.51	3.86	4.71	0.94	2.23	1./6	5 5	
38	36393	R25823	11.96	29.92	2.50	3.52	6.33	1.90	1.39	6.6 1.80	10.1	
585	460039	AA676740	2.76	6.89	2.50	1.35	1.80	1.09	4.19) ; ;		۰ ،
98	590148	AA156030	0.11	0.27	2.50	2.80	0.09	4.82	2.90	11.6	12:1	• ^
267	272600	N35922	1.96	4.89	2.50	2.26	3.56	3.63	1.29	70.7	0.1	۰,
88	345559	W73882	1.10	2.74	2.50	1.43	1.30	2.43	4.87	3.20	6. C	
3 8	1481228	AA83299	98.0	2.35	2.49	1.18	1.68	2.43	3.23	3.94	2.40	4 6
	1877733	A A D 4 2 5 8 2	2.47	6.13	2.48	1.32	2.39	3.08	3.67	2.69	1.75	ν (
	428626	AA703048	7.7	1.83	2.48	1.78	1.10	1.39	4.47	3.20	2.96	N (
	70000	A 4 8 0 8 3 0	9	<u> </u>	2.48	3.17	2.83	2.41	3.29	1.85	¥. ;	N (
7/0	180070	AA408038	3 6	- q	2.48	1.01	2.44	2.81	4.55	3.01	1.05	7
	130730	KZZU00	2 6	- 6	2.47	2.13	4.23	3.13	2.67	1.93	0.76	8 (
4/6	276.783	LZ8/LH	9. 0	24.33	2.47	2.30	3.42	2.22	3.78	1.28	1.8 <u>4</u>	~
2	70107	700 O	5	i > :	: i		17					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold of higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

					of Tumor			A17671101177	Ecidimon499/A	Fold(MPM123/A F	old(MPM125/A	# of Tumor
	IMAGE Clone ID	~ «	AveBPH	AveTumor	Fold(AveTumor/Av eBPH)	r Fold(MPM120/A	Fold(MPM121/A	veBPH)	veBPH)	VeBPH)	veBPH)	Samples
			6	,	. 247	3.23	5.03	0.73	2.49	2.35	1.00	7
576	209194	H63952	89.70 10.00	5.5	1.7	203	3.49	2.05	4.93	1.49	0.85	~
211	308466	N95495	5 . 6	50.81	2.48	3.20	5.99	1.70	2.59	0.95	0.35	7
578	785849	AA449120	50.0	0.0	2.48	1.18	1.24	4.12	5.65	0.95	1.63	7
2/6	158102	H52446)	2.48	<u> </u>	3.10	3.00	3.31	2.60	1.46	7
3	0/0897	N62/60	3.0	7 K3 L	2.48	7.96	3.28	<u>.</u> 2	96.0	0.82	0.71	8
8	8070	1/0436	30.0	3 2	2.48	1.10	3.69	3.77	1.95	2.40	1.86	74
8	19062	1400CX	8 t	50.02 4 20	2.45	2.15	1.50	2.68	1.02	3.10	4.26	81
3		AA151817) A C	9 07.	2.45	1.24	2.32	2.05	2.52	3.50	3.07	7
ž	745200	AA625864	2 6	<u> </u>	2.45	304	2.79	1.45	1.12	2.10	4.21	64
585	785890	AA449362	6.28	10.01 80.01	5.45 5.45		. E. 39	1.82	1.21	1.17	0.50	7
288	358046	W94591	19:0	94.0	C#.7		0.47	1.56	0.77	1.51	7.13	7
287	263860	AA101173	0.03	90 9	2.40	0.40	0.07		5,45	0.62	5.82	8
588	785407	AA453518	0.13	0.32	2.4	5. C			2.58	3.31	3.12	7
589	1460895	AA890098	5.95	14.53	2.4	2.08	6.10	62.0	1.92	2.27	2.80	8
280	753162	AA400457	2.39	5.8 2	2.44	3.25	3.02	9 6	10.1	58	1.88	8
591	712499	AA278382	2.63	6.41	2.44	1.50	3.82	0 4.4 0 4.0	i c	0.24	5.92	84
592	489656	AA099369	9.0	0.10	2.44	4.08	1.37	10.2	8 8		86.8	~
593	110503	T89996	0.60	1.45	2.43	0.73	1.24	0.78	0.92	0.00	4.47	
28	197465	H52048	0.35	0.85	2.43	1.53	2.49	3.94	18:0	47.4	÷ 6	
595	429729	AA011685	1.61	3.91	2.42	2.40	4.68	1.51	3.19	1.84	C.83	4 (
3	731371	AA421047	0.03	0.07	2.42	0.89	3.84	2.29	2.95	0.33	4.12	7 (
2 6	782585	4447527	048	1.17	2.42	3.45	1.49	5.47	1.78	. 4	0.80	74
	811010	AAABASAK	900	41.0	2.42	1.99	2.41	3.45	1.97	0.58	4.13	8 4
	742448	A A E O D 3 B E	2 2	7.78	2.42	2.19	3.40	1.76	1.31	2.53	3.33	~
	207328	No2470	800	0.07	2.42	5.00	3.80	0.67	1.74	2.14	1.15	~
3 5	207439	Wna687	24.42	29.00	2.42	2.70	4.38	0.99	4.78	1.4.	0.25	8
5 8	245320	NEZZEB	120	2.89	2.41	4.58	3.25	2.53	2.71	0.66	0.75	8
3 8	140057	D85083	0.75	1 82	2.41	1.19	2.77	2.50	3.00	1.88	3.13	α
3 8	433304	A A 600707	180	3.85	2.41	1.23	3.14	1.87	1.24	4.06	2.93	~
3 8	812227	AA455369	4.47	10.79	2.4	3.67	5.67	2.04	1.25	0.84	0.98	8
8	1580977	A4975010	1.07	2.57	2.41	2.29	2.67	3.01	3.69	1.52	1.27	7
202	283379	N52767	700	0.10	2.41	4.36	4.19	2.02	0.77	0.87	2.25	81 -
Ş	85195	771360	0.02	0.05	2.41	4.76	0.45	0.29	0.45	8.08	0.45	7
809	813310	AA455164	3.55	8.52	2.40	2.76	4.07	1.41	2.17	3.03	0.89	7
}	1						18					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

		Gen Bank		•		A/UCHAIMINATOU/A	Fold(MPM121/A	Fold(MPM124/A	Fold(MPM122/A	Fold(MPM125/A Fold(MPM121/A Fold(MPM124/A Fold(MPM122/A Fold(MPM123/A Fold(MPM125/A # of Tumor	old(MPM125/A	# of Tumor
	IMAGE	Accession	AveBPH	AveTumor	Fold(AVe i umon/A eBPH)	veBPH)	veBPH)	veBPH)	veBPH)	veBPH)	Verby)	Saldmas
		Number				1	6	0.85	1.86	4.32	1.76	- 7
ž	242037	H93328	0.38	0.89	2.33	3.07	2.13	3 3	233	3.03	1.50	8
645	470914	AA032084	13.04	30.41	2.33	2.36	3.53	1.24		5.53	4.59	8
878	38260	R49462	0.70	1.64	2.33	1.90	0.47	78.0		150	1.32	7
7.78	50250	H17800	0.81	1.89	2.32	3.30	3.08	2.73	2.02	 	0.82	8
878	743217	AA400128	1.85	4.30	2.32	1.78	3.31	2.05	2.70		271	. ~
3 3	200844	N74082	25.	10.65	2.32	3.15	3.16	0.87	2.32	2 !	. ç	
Š	10007	700000	2	13.75	2.32	1.14	2.10	1.47	3.01	79.4	9 1	
650	814585	AA480906	b i	5	33	1.73	3.33	3.46	2.02	1.79	36.	۷ (
651	240766	H80215	£.74 0.01	0.50	10. c	1.25	1.34	1.59	3.79	2.55	3.37	N (
652	34869	R44447	2.35	0.40	 	4 10	1.12	1.78	1.85	1.37	3.63	7
653	51263	H18838	0.0g	0.14	F.2.	5 4	96	5.47	1.37	3.40	1.28	01
3 2	810209	AA464522	1.10	2.54	2.30	J.35	9 6	3 20	5,16	0.46	0.83	7
655	795831	AA461508	0.07	0.17	2.30	1.39	85.2 20.0		2.67	1.74	0.77	7
656	745138	AA626698	12.76	29.36	2.30	3.25	2.34	t 6	247	0.55	0.48	7
657	489076	AA057195	0.05	0.05	2.30	1.88	3.30	2.10	181	0.98	1.43	7
658	344108	W73781	3.42	7.83	2.29	4.00	3.17	7:30	5 6	3.58	1,55	8
659	727192	AA283819	12.78	29.26	2.29	2.09	0.89	2.24	- t .	7 40	1.60	8
088	41541	DK2780	4.22	9.64	2.29	2.19	3.57	3.59	1.20	e :	700	•
3 8	705353	A A 4 53487	18.3	9.86	2.29	1.68	4.15	1.50	4.56	9.0	7 20	
8	2000	1010010		0 71	2.28	2.41	0.86	1.19	0.31	4.58	9.40	4 (
995	40400	K52088	- t	- 6	2 28	1.85	3.18	2.62	2.50	3.07	0.48	7 (
883	48330	H14949). (0.38	07.7	3 18	0.12	2.08	2.75	5.09	0.47	м
8 8	769600	AA425900	0.09	0.20	2.28	2 9	. e	3.29	1.78	1.99	1.51	7
665	112865	T87139	2.61	5.95	2.28	8C.1	9 6	8 0	3.14	0.18	2.05	8
888	362795	AA018683	90.0	0.13	2.28	2.04	0 0		1 96	3.17	1.91	~
687	384397	AA708676	0.46	1.05	2.27	1.47	3.56	t :	2 6	1.53	5.05	8
868	26568	R39111	0.74	1.68	2.27	3.15	0.55	0.70	5 5		1.65	2
689	199228	R95811	0.58	1.27	2.27	1.69	3.64	3.07	 	5 4	4.27	8
07.0	41070	DAR400	0.03	0.07	2.27	1.48	2.18	0.33	2 :	<u> </u>	į	
2 6	K4403	140047	0.37	0.83	2.27	0.74	4.79	4.4	1.55	3.13	<u> </u>	
5	2	1176111		2	7.00	3.52	2.57	3.03	2.05	D.61	· ·	. (
672	346552	W74377	70.0	\$ 6 5 6	 80 C	3.02	3.29	1.52	2.48	1.77	1.51	N
673	624785	AAT8T885	5	9.02	9 00	88	3.59	9. 9.	2.27	1.42	1.67	N
874	626716	AA191548	17.62	39.85	07:7	9 9	3.40	2.74	2.58	0.71	0.53	7
675	785760	AA449686	0.02	0.0 4	2.26	5.50 1.00	6 CY	. 16	1.92	0.60	1.62	7
878	345145	W72231	3.57	8.07	2.26	3.73	3.32	2 6	1.49	3.82	4.13	~
677	46620	H10036	15.87	35.82	2.28	0.96	7.07	70:1	<u>:</u>			
							20					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold of higher in at least one fumor sample or 3 fold or higher in at least 2 tumor samples.

	IMAGE	Gen Bank Accession	AveBPH	AveTumor	Fold(AveTumor/Av Fold(MPM120/A Fold(MPM121/A Fold(MPM124/A Fold(MPM122/A Fold(MPM123/A Fold(MPM126/A # of Tumor earth very) very) very)	old(MPM120/A	Fold(MPM121/A veBPH)	Fold(MPM124/A veBPH)	Fold(MPM122/A veBPH)	Fold(MPM123/A veBPH)	Fold(MPM125/A veBPH)	Samples
	Clone 10	Number			ì			23	0.21	5.03	4.87	8
878	811832	AA463627	9.02	0.11	2.25	0.87	17.0	2 7	17	1.38	3.78	N
67.0	49141	H16554	0.19	4.0	2.25	1.32	2.11	<u>.</u>	 8.	1.62	2.28	М
	822804	A A 4 8 0 7 5 2	2.53	5.67	2.24	3.20	3.06	5 .	200	0.37	0.36	7
3 3	364437	A 4 4 7 8 8 2 7	35.6	7.92	2.23	2.21	5.87	0.66	r r	77	1.75	8
ē 8	131401	A A 8 8 2 8 0 0	9	+	2.23	1.70	1 .3	96.0	7.5	; ;	1.63	8
200	80010¥1	AA663600	3 6	1	2 2 2	1.86	4.17	1.42	3.53	4.5		•
683	45089	H05140	0.32	2 5		3.16	3.03	<u>4</u> .	1.7	1.67	7/1	4 (
2 8	232912	H72643	21.80	48.25	7.7	200	3.06	2.59	3.48	0.79	1.35	7
685	135789	R33353	0.11	0.25	2.21	/8'1	3 33	1.18	2.96	1.72	0.26	8
888	795308	AA454149	0.10	0.23	2.21	3.92	3.62	3.33	2.25	1.21	3.10	7
687	448270	AA777696	90.0	0.13	2.21	2.46		6.6	2.18	2.77	3.05	8
888	808108	AA698587	1.71	3.77	2.20	1.65	3.01	BC.U) i	18.0	3.26	8
3 8	124073	B23241	4.56	10.05	2.20	1.61	1.24	1.02	. i	108	0.90	8
9 9		70724	8	2.18	2.20	1.77	3.15	3.31	01.2	3 4	3.27	7
250	733711	+R+//L	2	2 6		2.38	1.42	1.49	3.06	RC.	4.0	
691	243800	N39338	1.48	3.27	0.20	1 26	3.11	6.76	0.80	0.20	1.07	7 (
692	782541	AA448486	0.13	0.29	2.20	03.1		1.19	1.36	3.30	3.35	74
693	813641	AA447744	0.48	1.06	2.20	49.		3.24	1.57	1.12	1.09	84
469	126568	R06862	3.87	8.47	2.19	2.26	3.00 10.00	, c	0.20	0.20	5.50	7
695	345553	W73889	0.05	0.11	2.19	1.87	1./3	9. 6	3.42	3.28	0.98	7
888	51947	H24313	2.05	4.49	2.19	1.10	1.74	7.01	; e	1.83	6.60	7
709	40010	B54038	0.09	0.21	2.19	0.84	0.11	3.51	2 6	4.87	0.81	7
	183188	A A 0.74224	0.29	0.64	2.18	1.56	3.14	2.20	0.53	, t	2.76	74
	2000	75030	30.0	0 11	2.18	3.98	1.50	0.27	3.22		103	8
20 1	15557	RCOCC I	300	8	2.18	0.48	0.82	6.02	3.04	0.70		۰ ۱
8	188070	AA418/48	3	200	2 43	2.11	3.23	3.65	1.29	1.24	<u> </u>	
2	33860	R44840		15.50	2 4 6	4.03	4.62	2.45	1.01	0.17	0.77	4 (
702	46166	H08018	5.56	8 6		3 17	3.06	0.65	2.60	0.77	R/ 7	4 (
703	283191	N51362	2.72	08.6		1 43	3.18	0.73	2.68	3.94	- 8	NI (
ş	647985	AA207105	-	24.76	71.7	G 6	28.0	1.27	1.28	4.03	4.23	N
705	491001	AA138710	5,71	12.38	2.17	97.1		3.92	3.45	1.19	1.39	7
92	1481161	AA868038	1.08	2.34	2.18	1.6/	2	1 40	3.60	1.13	0.68	81
707	757503	AA426380	1.95	4.21	2.16	1.68	 	9	900	0.72	1.23	8
200	280375	N47113	1.16	2.51	2.16	4.65	3.70	0.5) ()	80 0	0.92	7
9 6	383918	AA021247	1.43	3.09	2.16	1.68	3.74	1.8.	9.00		3.09	8
	454720	T0800+	0.73	1.58	2.15	1.34	3.07	1:1	1.30	6.30 0.00	3 20	2
2 ;	8211.21 828804	AADADORD	1 29	2.77	2.15	1.58	0.79	0.98	5.28	90.1	2	1
=	3	ANDROS	<u>:</u>				21					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

		3			;	4000	Cold/MDM424/A	Fold/MPM124/A	Fold(MPM122/A	ESILIMBA121/A FOIGINPA122/A FOIGINPA122/A FOIGINPA123/A FOIGINPA123/A FOIGINPA123/A FOIGINPA123/A FOIGINPA123/A	old(MPM125/A	# of Tumor
	IMAGE	Accession	AveBPH	AveBPH AveTumor	Fold(AveTumor/A eBPH)	(V FOIG(MFM120/A VeBPH)	veBPH)	VeBPH)	VeBPH)	veBPH)	veBPH)	Samples
		Number				•	60	1 33	3.92	1.95	3.18	-
712	207370	H58834	0.13	0.27	2.15	1.14	05.1	8:-	700	1.88	0.96	7
713	202795	H53964	2.15	4.60	2.14	1,36	3.31	3.00	1 1 1	76	48.0	8
	828138	AA521337	0.0	0.02	2.14	4.28	0.94	4.27	Q 7		1.13	7
	3224R1	WHENDE	2.11	4.50	2.13	3.57	3.32	2.78	V.0	5 4	7 7	
2 4	188232	H45711	0.61	1.31	2.13	3.34	1.13	1.3 4	1.68	2.10	0.47	. 6
2 ;	10000	10000	2.60	45.5	2.13	2.40	3.78	3.35	2.45	t (5,0	080	. ~
	C7C / & I	100701			2 13	2.50	3.41	1.76	3.41	20. L	80.	
718	460279	AA677629	9 9	ţ ;	; ;	1 10	2.23	1.40	3.97	3.07	1.03	v
719	245444	N53492	0.53	1.12	2.13	000	3.50	3.33	2.50	0.59	1.88	7
720	83029	T67807	1.42	3.02	2.13	9 6	9 6	1.08	0.19	1.11	3.34	74
724	276871	N39426	0.05	0.11	2.13	9.80		200	85. 46.	1.68	0.08	~
227	417059	W87801	0.12	0.25	2.13	0.40	t €		123	1.67	2.04	8
723	294255	N99839	13.11	27.87	2.13	1.47	3.12	5.45 7.89	9 5	4.42	0.94	8
724	243700	N45083	1.34	2.84	2.13	1.35	3.02	7.07 7.07	 8.	091	2.84	7
725	285040	N63375	5.95	12.63	2.12	3.18	3.00	G !,	07: 4	0.82	1.02	7
728	73252	T56013	1.33	2.82	2.12	1.70	3.05	1.47		000	5.71	7
2	743465	AA609385	1.79	3.78	2.11	0:00	1.15	3.08	78.0	1.40	1.27	8
Ş	464486	DETABLO	F 3	80.6	2.11	3.15	3.51	2.17	01.1	o e c	5	. 6
9 6	20170	A 4 E E 2 2 7	5 0	0.27	2.10	3.09	96.0	3.49	2.09	0.80	5.5	۰ ،
87/	00/4/0	1776C+WW	2 2	i 6	2.10	1.29	1.12	3.41	1.85	1.43	n e	• ‹
8	322005	W37418	5 6	5 6	; ¢	19	3.31	3.19	1.59	1.17	1.68	,
73	416945	W87710	3.08	6.40	2:10		3.14	1.32	2.68	3.20	1.03	N
732	624429	AA182680	6.29	13.17	2.10	1.20		0.74	4.67	1.56	4.35	7
733	1031844	AA609686	0.08	0.16	2.09	0.17	9. 3		4.29	4.03	0.28	7
ş	345176	W72263	0.18	0.37	2.09	1.24	1.24	2 ·	3.65	0.99	1.11	7
735	878652	AA670200	1.48	3.08	2.09	- - - - - - -	3.20	5 6	1 55	1.49	2.04	8
736	366511	AA026609	4.79	8.99	2.09	1.37	3.02	000	0.87	1.39	1.95	7
737	235173	H73013	<u>+</u> .	2.99	2.08	1.48	3.27	000	3 73	4.93	0.84	7
738	296448	N74623	90.0	0.13	2.08	2.84	0.16	0.50	4 6	38	44.0	8
730	357284	W03585	1.1	2.28	2.06	2.07	3.92	0.82	3.60	9000	. A	8
		000000	2	79.0	2.05	1.08	3.65	1.82	0.25	0.40		
? ;	RAVAS	N32040	3 5	2.45	207	0.92	3.31	1.69	1.59	3.62	+ :- (4 (
•	/CR70+	C74700W	2 6	, t	70.	4.41	2.53	0.14	0.63	1.16	3.40	4 (
742	282770	N63646))	<u>.</u>	3	3.70	3.56	1.95	1.90	0.32	0.80	7
73	195635	R89317	0.26	50.0	t 6	0.10	3.21	3.13	2.60	9.0	0.50	7
7	583873	AA101299	0.58	1.19	2.03	2.3	0.00	0.98	3.23	5.47	1.68	~
745	795522	AA454218	0.05	0.10	2.03	00.0		•				
							•					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one furnor samples.

		Jue 0			1	A/00/18/00/19/20/A	Eold/MPM121/A	Fold/MPM124/A	Fold(MPM122/A		Id(MPM125/A	# of Tumor
•	IMAGE	_	AveBPH	AveTumor	fold(AveTumor/A eBPH)	V FOIG(MFM120/A VeBPH)	veBPH)	VeBPH)	veBPH)	veBPH)	veBPH)	Samples
		Number				,	4 4 5	88	1.13	3.33	1.27	7
746	42035	R59068	0.19	0.39	2.03	1.37	77:		780	0.40	4.35	8
747	280412	N47208	0.05	0.05	2.03	4.97	3.09	i i		1.37	3.99	2
748	700302	AA283693	9.0 40.0	0.09	2.03	1.01	0.21	27.5	S 4		3.88	7
749	40082	R52530	0.33	0.68	2.02	3.25	0.81	1.70	5.5	133	0.97	7
750	1486109	AA936783	112.27	226.02	2.04	1.69	3.11	00:1	1.12	2.10	3.08	8
751	415906	W86196	0.27	0.55	2.01	1.00	3.06	00.	3 47	, K.	2.71	8
752	399421	AA733177	2.08	4.18	2.01	0.73	0.73	00:- 6		1.55	0.99	8
753	744962	AA625907	10.80	21.68	2.01	1.77	1.69	0 4		1.36	0.77	~
\$	415111	W93147	0.65	1.30	2.01	1.54	3.53	10.5	<u> </u>	0.90	4.37	8
755	811046	AA485427	0.08	0.16	2.01	1.48	0.12	21.1	5 6	2.31	3.41	7
758	590640	AA158035	0.14	0.27	2.01	0.92	3.86	6.93 6.93	. 6.0	7.57	0.70	8
757	588053	AA135001	2.74	5.47	1.99	0.61	0.28	1.31	0.6	14.	0.87	~ ~1
758	278496	N66135	2.33	4.63	1.99	1.28	3.38	1.63	9.6	3.70	0.67	8
759	32962	R43910	0.33	0.65	1.98	0.56	4.06	88.	0.87	134	0.49	8
780	878785	AA670411	13.20	26.15	1.98	1.66	3.55	4.1	5.75 6.00	10.4	0.49	8
781	627105	AA190634	9.88	19.43	1.97	1.41	3.00	1.85	9. F. C	<u> </u>	0.42	~
782	42123	R59621	1.76	3.45	1.96	2.41	3.44	0.45	5.07	5 6	1 22	7
, E	245547	N55167	2.57	5.02	1.95	3.53	4.14	1.49	16.0	* 0	3.50	۰ ،
3 3	769704	0 0 4 4 7 7 0 3	0 13	0.25	1.95	1.25	3.84	1.08	1.57	0.60	9 4	
\$;	£0.70	7611187	2 2	882	1.95	0.96	3.61	1.32	3.38	1.26	81.T	4 6
62	17804	0770LH		9 6	1 95	3.04	3.06	0.98	1.89	1.68	5 .	7 1
8	1031309	AA609088	[6.5]	0.88	3 7	0.15	3.60	2.65	0.63	0.15	4.47	74
787	376043	AA040265	0:0	2 0	5 6	2.30	1.24	3.70	3.77	0.56	0.07	7
88	51373	H24006	0.13	0.20	<u>.</u>	7 2	0.87	1.45	1.35	3.22	3.15	7
92	361363	AA017706	20.0	10.05	 	0.78	1.30	1.49	1.78	3.10	3.1	6
20	121124	AA704537	R (2 6		- F	1.88	0.78	0.75	3.17	3.66	8
E	769933	AA430506	70.0	0.89	 	08.0	0.16	1.67	0.16	3.99	4.78	~
112	782826	AA448257	g (7	60.	4 97	0.83	0.83	0.96	0.89	3.01	74
713	841281	AA487203	70.1	28.7	76.7	90.	3.63	1.18	3.76	0.32	0.68	~
7	610883	AA172188	25. 28.	5.46	1.82	08:-	60 E	0.62	0.89	1.23	2.43	7
118	813730	AA453783	2.20	4.23	1.82	07:0	0.0	4.59	5.15	0.38	0.39	7
778	810896	AA485351	0.03	0.05	1.92	0.90)) (7	60 40 60	0.92	3.71	7
E	841666	AA487561	5.91	11.33	1.92	1.22	60.00	÷ ÷	3.17	0.78	40.5	84
778	277112	N39611	0.73	1.38	1.91	2.14	3.20	7	. ¥6	88.69	5.59	7
417	530608	AA071089	0.05	0.10	1.90	0.18	9.18	¢.	3	}		
							23					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor samples.

		Gen Bank			Fold(AveTumor/Av Fold(MPM120/A Fold(MPM124/A Fold(MPM122/A Fold(MPM123/A Fold(MPM126/A # of Tumor	Fold(MPM120/A	Fold(MPM121/A	Fold(MPM124/A	Fold(MPM122/A	Fold(MPM123/A	Fold(MPM125// veBPH)	A # of Tumor Samples
	IMAGE	Accession	AveBPH	AveTumor	eBPH)	veBPH)	veBPH)	Vebru)			**	•
	Cione	Number				7	0.48	2.48	3.11	3.59	<u>*</u> :	
780	208570	H63223	0.59	1.10	1.89			1.57	0.87	<u>5</u> .	3.49	7 (
781	131887	R25166	2.12	3.98	1.88	3.19		8	1.31	0.70	1.31	8
,	241078	W81361	1.28	2.41	1.88	3.07	3.00	- č	3.24	0.89	0.92	8
1 0	484727	AA682283	1.49	2.80	1.88	3.10	0.77	£ 2.5	0.50	4.16	3.30	8
3 3	945550	A4456818	3.03	5.67	1.87	0.83	0.89	ţ 6	4.56	0.80	0.48	7
5 5	33000	B44538	7.22	13.49	1.87	0.98	3.01	1.38	120	0.08	3.55	7
3 8	34045	R44754	0.12	0.23	1.87	0.95	3.90		55.55	0.24	0.23	М
3 6	743189	44401434	0.0	0.08	1.86	2.05	3.30	0.7	3.14	1.38	1.06	8
4 6	247194	N54036	0.36	99.0	1.86	1.09	3.23		3.07	1.82	0.79	8
8 8	183771	14448	1.91	3.56	1.86	1.23	3.04	77.1	1.47	1.48	0.13	7
9 6	491764	AA150502	0.08	0.14	1.86	3.80	3.09	07.1	109	0.42	0.42	74
į	782504	AA447531	0.02	5 0.0	1.85	3.26	3.78	. i	0.47	1.55	0.39	7
6 6	471859	AA035144	0.03	0.05	1.85	3.10	5.05	. c	1.75	0.10	3.28	7
2 6	278712	N34895	0.13	0.23	1.85	0:30	0.39	9.70	3.22	1.20	0.59	8
3 3	77487	H08154	1.53	2.83	1.85	2.02	3.26	8 70 T	88 8	0.39	0.42	7
\$ \$	284274	1000	2.71	2.00	1.84	1.16	3.49	D. 10.	6.6	0.60	0.98	7
	830084	A 487508	11.79	21.75	1.84	3.03	3.73	0.71	. 89	0.19	4.08	7
8 5	608001 E00808	AA124578	0.88	1.61	1.84	4.66	0.80	0.64	3.24	1.13	0.68	7
200	77877	158937	0.29	0.52	1.83	1.41	3.32	E 7	+ 64	3.17	0.53	7
1 2	830820	44489782	0.23	0.41	1.82	1.28	3.14	1.17	22.0	4.18	3.24	7
B 8	K3004	R15784	0.11	0.20	1.82	1.57	0.53	0.62 4.45	0.85	0.69	3.60	7
3 5	258531	W96134	2.78	5.05	1.82	3.97	0.53	5.5	3 10	0.42	2.00	7
5 6	32339	R42823	0.14	0.24	1.80	0.81	0.47	0.50 M	103	3.31	4.34	7
8	417688	W89074	0.20	0.35	1.80	1.48	0.57	6.50	1.30	0.25	4.72	8
ğ	1435339	_	9.0	0.07	1.79	0.22	6.04 6.04	9 66	1.30	0.20	0.43	63
805	128973		0.05	0.09	1.75	0.93	77.4	3.73	3.19	0.89	0.86	N
808	361323	AA017544	2.04	3.57	1.75	0. 48.0	9.7	17.	3.4	0.20	4.04	01
807	586440	AA148862	0.05	0.09	1.75	0.31	0.7	96.0	0.48	0.90	3.53	~
808	857196	AA773139	0.46	0.79	1.74	1.59	3.02	3.02	0.31	0.31	0.31	~
8	586958	AA130984	0.03	90.0	1.71	6.02	6.0	0.62	0.74	3.11	3.35	8
810	741891	AA401972	5.81	9.8 78.	1.69	1.65	80.0 C80	76.0	3.45	0.57	0.18	~
21	810027	AA455275	0.0	0.07	1.69	6.23	0.33	0.21	3.08	0.21	1.05	7
812	811609	AA455020	0.03	0.02	1.65	5.05 0.45	3.01	1.77	0.79	0.33	3.45	Ν.
813	361807	W92431	0.21	0.35	1.63	î S						
							74					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold of higher in at least one turnor samples.

		Gen Benk				Fold(MPM124/A Fold(MPM121/A Fold(MPM124/A Fold(MPM122/A Fold(MPM123/A Fold(MPM123/A Fold(MPM126/A # of Tumor	Fold(MPM121/A	Fold(MPM124/A	Fold(MPM122/A	Fold(MPM123/A	Fold(MPM125/A	# of Tumor
	IMAGE	•	AveBPH	AveBPH AveTumor	roid(Averdinon) eBPH)	veBPH)	veBPH)	veBPH)	veBPH)	VeBPH)	(Luage)	
		Number			•		6	0.51	3.09	3.13	1.83	64
814	795538	AA459651	0.10	0.16	1.60	0.95	0.10	5 6	636	0.68	3.79	7
	280488	N32071	0.02	0.02	1.59	0.51	99.0	0.38	700	67.0	0.43	7
2 6	2000	1 10201	0	700	1.59	0.28	4.10	0.43	3	? ·	4.53	~
818	CSAROS	AA168134	3 6	5 6	1 87	0.21	3.55	3.54	0.45	0.10	70.1	
817	47187	110404	0.00	0.0		5	3.48	3.31	9.0	0.18		7 (
818	810948	AA459383	0.08	0.09	/0.1	9 6	0.23	1.13	1.63	3.05	0.27	74
819	490556	AA100696	9.0	0.07	1.58	3.03	9 6	3.62	0.20	0.18	0.34	8
820	769579	AA425826	0.11	0.17	1.53	4.53	0.38		151	0.22	0.22	7
821	454083	AA676998	0.05	0.07	1.47	3.29	3.40	77.0		0.31	0.67	7
2	52071	H24350	0.27	0.39	1.47	3.02	3.31	69.0	5 6	33	0.33	~
2	811066	AA100298	0.03	9.0	1,35	0.33	0.48	3.30	t ::	0.40	3.07	~
2	742818	AA400482	0.24	0.32	1.33	3.06	0.05	4	0.23	9 4	0.17	-
3 6	E04484	AA462478	90.0	1.88	32.02	186.97	2.01	0.17	7.01	15.6	232	-
200		NO SECON	200	1.85	26.82	2.37	0.45	0.1 4	41.0	100.4	2	•
979	2000	88078N	2 6	80.73	22.47	0.60	1.38	132.15	0.22	t i	10.0	• •
827	4523/4	AA/008/8	6.0		16.48	2.18	2.14	2.48	0.75	91.20	t (- ,
828	384015	AA702640	0.32	97.6	2) (i	97.0	0.29	2.14	46.98	2.25	- -
829	46641	H10344	0.0	9.0	8.77	0.00	7.0	1.78	0.14	40.15	1.39	-
830	158322	R72618	0.08	0.61	7.60	1.5/	9 6		15.	0.58	33.43	-
831	81413	T60161	0.15	0.97	6.43	0.62	58.0	2	2	2.90	2.23	-
833	745232	AA626152	0.33	1.99	6.03	2.98	2.15	80.47	- +	20.0	31.39	-
3	420240	A A 0.07.278	0.27	1.55	5.77	0.90	0.45	0.30	1.33	9 6	28.0	-
3	017874	0.21000	900	1 48	18.5	0.07	1.18	0.70	30.40	80.0	8 5	. •
*	50354	H17804	0.40) E (0.52	0.99	0.13	1.21	1.70	RR: 17	- •
835	364563	AA022949	0.30	1.65	3.42	10.0	68.0	0.63	1.19	1.10	27.41	-
8 38	488422	AA044862	0.05	0.27	5.40	2 6	5 7	2.14	1.43	1.89	24.70	~
837	50789	H17630	0.30	1.59	5.36	0.82	17:	181	1.51	24.08	1.74	-
838	302180	N79989	0.22	1.16	5.35	2.39	n (0.28	97.	0.28	-
839	128083	R09747	0.04	0.20	5.12	1.88	30.1	01.07	1.48	0.14	0.14	-
840	25933	R36969	0.07	0.37	4.96	0.96	0.82	70.20) to	800	0.43	-
77	229ABB	HBB542	0.11	0.53	4.77	2.48	0.09	25.48	0.07	3 .	0.62	-
5 3		10001		187	4.71	0.95	1.14	22.95	1.24	0.70	9 6	• •
7	144021	AA621310	4 4	9 6	4.42	0.13	1.05	0.32	2.09	22.55	0.40	- ,
8	884462	AA629707	cr.o	0.00	*		1.15	1.35	20.54	0.78	2.08	-
3	361291	AA017468	9.	14.4	4.41	/6:0	- 6	1 98	0.42	1.17	0.09	- -
25	301735	N90882	0.11	0.49	4.33	19.43	2.09 2.09	 85. cc	0.22	0.21	0.41	-
878	753076	AA436479	2.41	10.37	4.30	0.47	69.0	63.63	ļ č	69.0	2.52	-
847	510381	AA055585	0.32	1.35	4.23	19.31	0.51	1.32	<u>.</u>	;		
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Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

		Gen Bank			ESIMANSTIMENTAN FOLDIMPM120/A FOLDIMPM124/A FOLDIMPM124/A FOLDIMPM122/A FOLDIMPM123/A FOLDIMPM123/A FOLDIMPM125/A # OF Tumor	Fold/MPM120/A	Fold(MPM121/A	Fold(MPM124/A	Fold(MPM122/A	Fold(MPM123/A	Fold(MPM125/A	* of Tumor
	Clone ID	Accession	AveBPH	AveTumor	eBPH)	veBPH)	veBPH)	veBPH)	veBPH)	veBPH)	VeBPH)	Samples
•	,	Jegmun.	37.0	6	7.54	0.91	1.74	18.40	0.98	2.09	1.12	-
3	44840	AA///536		- 6	4.4	1.19	1.74	0.82	1.23	19.01	0.85	-
2	504940	AA150619) (2 ;) F	1.32	2.06	1.18	1.28	2.94	15.68	₩.
200	287771	861881	, c	+	SO:+ EO ₹	1.32	1.09	0.51	0.38	20.03	0.86	-
2	74107	**********		3 6	3.97	1.74	1.54	0.57	0.04	17.96	1.99	-
200	1 0000	AA034014	2.0	30.0	3.82	2.13	2.98	1.30	1.23	14.18	1.67	-
2	17467	K30201	9	7 F. F.	3.80	1.08	1.29	15.59	1.72	2.23	0.91	-
8	117470	A4480875	2 5	2 6	3.74	1.37	0.95	2.14	0.67	14.52	2.78	~
2 2	43065	761886 8868	7 7 0	. t.	30.00	0.59	0.93	1.22	1.92	1.22	15.62	
8	24/103	W80000	9 6		3.48	2.06	1.14	1.62	10.80	2.71	2.56	4-
824	32381	R43452) o	5.5	6. 6. 6. 6.	0.47	0.21	1.36	0.63	1.95	16.16	-
8	2//ZAC	112821AA	t 6	9 9	3.45	0.77	0.38	1.22	14.94	1.32	2.08	
	008404	A401/403	3 6	9 6	344	1.52	1.27	1.69	15.08	0.51	0.39	~ -
3	340041	M26/60	7 6	900	340	16.70	0.81	0.76	0.95	1.03	0.13	-
5 8	77//RQ	AADBABAS	9 6	9.60	338	17.69	1.01	0.60	0.60	0.21	0.05	_
700	700000	AA018332	. e	222	3.35	1,17	0.97	14.33	1.21	1.19	1.22	.
3 3	150004	A A GORFOR	27	56	3.34	1.80	1.58	1.72	1.32	11.93	1.68	-
8 8	788280	AA424050	2 6	2.69	3.28	1.66	12.82	2.38	0.89	1.08	0.83	- .
2 8	46030	M40028	0.28	0.92	3.27	1.15	0.99	0.74	1.64	13.18	4.89	.
3 2	1035765	AAR29117	10.22	33,26	3.25	0.55	0.26	1.17	15.74	1.49	0.32	- - ,
2	591025	AA159068	0.21	0.67	3.24	1.63	0.50	2.32	1.36	2.40	11.24	- ,
3 2	194515	R86242	0.58	1.87	3.24	6.47	2.02	2.78	2.94	2.78	2.43	- ,
870	308412	N93790	0.02	0.05	3.21	1.60	0.62	1.66	0.62	12.88	1.87	- ,
871	767449	AA418000	2.79	8.93	3.20	1.31	1.70	0.78	2.24	10.28	2.91	
872	809981	AA455197	0.0	0.28	3.19	0.83	0.72	0.87	9 .0	0.35	15.41	- ,
873	322175	W37778	1.80	5.71	3.18	2.63	6.05	2.60	2.45	2.92	2.41	- •
874	287770	N62251	0.10	0.31	3.14	1.79	0.93	12.98	2.84	0.10	0.23	- ,
875	51216	H18471	0.33	1.03	3.13	1.22	1.79	1.32	1.82	11.65	1.01	- •
876	591683	AA147214	0.03	0.10	3.13	0.62	0.57	11.65	2.53	1.55	 85.	- •
877	51951	H24317	0.33	1.01	3.10	1.50	12.98	1.31	1.06	0.77	00.1	- •
878	429175	AA004824	0.19	0.58	3.08	1.55	6.34	2.94	2.52	2.46	2.64	- •
879	3448	R44164	0.46	1.41	3.05	0.95	11.69	0.77	1.45	1.05	2.37	·• •
88	280217	N47941	6.29	19.12	3.04	1.96	1.84	0.85	0.83	11.68	1.12	- (
. 58	360838	AA017141	0.34	1.01	3.00	2.60	1.86	2.57	1.68	1 .18	8.11	_
							90					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold of higher in at least one tumor samples.

			-		TILLING ENGINEMASAIA FOLDIMPMISZIA FOLDIMPMI		A/4088494/A	Fold/MPM124/A	Fold(MPM122/A	Fold(MPM123/A F	old(MPM125/A	# of Tumor
	MAGE	Gen Bank Accession	AveBPH	AveTumor	Fold(AveTumor/A	V FOID(MPM120/A VeBPH)	veBPH)	veBPH)	veBPH)	veBPH)	VeBPH)	Samples
	Cione				•		6	1000	1.25	1.05	0.81	+-
883	730946	AA416543	1.1	3.32	2.98	0.80	0.96	0.71		121	0.81	-
2	706307	AA454155	2.01	5.93	2.85	09:0	0.67	13.6/	t ;		1.65	Ψ
3 3	400237	D95773	0.39	1.13	2.91	2.12	2.71	1.68	I • ./	.03	1,73	
	1000001	2710004	90	1.73	2.91	1.15	2.19	2.85	1.86	10.7		•
	10204	AA270445	304	8.74	2.90	2.81	2.95	1.84	5.15	7.00	40.04	•
	10407	2418134		1.82	2.90	1.39	1.34	1.32	1.43	9. 6	0 50	•
è	/2010/	1000000	3 6	2	2.89	0.52	1.94	0.52	2.68	81.7	40.0	
88	625516	AA18/595	9 9	3 7	2.88	1.28	0.45	1.22	10.05	<u> </u>	2.63	- •
88	838478	AA457517	9. ;	97.7	3 c	2.57	7.47	2.29	1.27	1.75	T8.1	- ,
880	46367	H08828	-	90.4	00.7		2.40	2.53	1.74	7.18	1.79	_
891	213753	H72322	0.28	0.73	2.87	1.07	2.25	1.77	1.73	9.16	1.31	-
885	126449	R06706	0.22	0.63	2.87	/a:0	3 4	80.0	1.18	2.07	6.39	-
893	305243	N95007	09.0	1.70	2.86	1.99	2.55	20.00	103	1.03	1.16	-
36	254117	N22230	1.22	3.49	2.86	0.86	0.81	47.71	. c	7.24	1.48	-
895	199285	R95893	0.14	0.41	2.85	2.05	2.36	08.1		, Z	1.98	-
8	51918	H22563	9.74	2.10	2.84	2.50	2.20	2.20	70.7	2 87	7.65	-
	444370	T86173	1 89	, se	2.83	1.63	1.87	1.92	5.	60.7	30	-
	376434	A 5020743	1 2	334	2.83	1.76	2.11	0.92	20.7	60 F	20.6	. +-
22	0.00/0	AA0387 13	2 5		2.81	1.10	2.13	1.85	1.95	7.78	70.7	- 🕶
688	24676	. 438808	9 6			121	0.97	10.42	1.72	1.24	ר קי ן	- ,
8	127931	R08938	0.66	 	87.7		0.49	0.62	2.18	0.57	11.87	-
8	77915	T61323	8.10	22.59	R)'7	90.0	90 6	2.59	5.69	2.57	1.69	-
805	327461	W20438	0.76	2.12	2.78	2.03	50.4 00.00	1.45	1.03	2.33	1.61	-
903	40283	R55261	0.15	0.41	2.78	48.1	0.29	283	1.66	2.95	5.61	-
Š	826109	AA521327	0.40	1.	2.78	1.65)) i	1.30	1.08	1.26	₩-
905	113160	T83842	99.0	1.83	2.77	1.55	S	- 6	99	10.48	0.03	
8	288768	N62498	0.35	0.96	2.77	1.16	00:1	2 6	8 37	1.28	1.29	-
200	289936	N59336	0.57	1.58	2.76	1.73	2.11	00:1		0.69	1.02	-
Š	770868	AA434487	0.09	0.26	2.75	12.49	0.46	0.72	? ;	3	1.48	-
8	197779	R93515	1.43	3.91	2.74	0.94	1.29	10.19	L.;	80 80	1.62	-
3 3	70007	0.0000	70	2.57	2.73	1.07	1.33	9.71	1.37	07:	5 5	• •
916	140831	K6/218		7 7 7	2.72	1.73	1.78	2.68	6.30	1.99	1.87	- •
<u></u>	84/1C	HZ4328	8 6		77.6	1.62	1.49	1.30	10.18	0.79	08:0	- ,
912	280977	N50864	3.			8 20	1.63	1.75	1.84	1.63	1.16	_
913	41835	R54212	0.33) (A)	2. c	33.	2.72	2.53	6.58	1.45	0.60	-
914	951125	AA620556	15.08	40.69	2.70		1.50	2.02	7.45	2.53	1.28	~
912	51011	H19242	4.02	10.78	2.68	- 2	1					
) (

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold of higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

		1100			:	Toldimph122/A Foldimph122/A Foldimph122/A Foldimph122/A Foldimph123/A Foldimph126/A # of Tumor	A/1-0-14/19/04/1/	EOIG (MPM124/A	Fold(MPM122/A	Fold(MPM123/A F	old(MPM125/A	# of Tumor
	IMAGE	Accession	AveBPH	AveBPH AveTumor	Fold(AveTumor, eBPH)	/Av. Fold(MPM12U/A veBPH)	veBPH)	veBPH)	veBPH)	VeBPH)	veBPH)	Samples
		Number				12	2.81	2	1.80	5.50	2.76	-
916	44303	H06380	1.99	5.32	2.68) (C.)	- C	2.29	1.39	1.88	2.19	-
917	296838	W01211	0.61	1.63	2.68	1.80	4 6	1 +	2.15	1.48	1.91	-
918	139641	R64231	0.12	0.31	2.67	36.1	7.58	2 6	1.38	0.33	10.39	-
918	609228	AA179161	0.09	0.25	2.67	0.51	0 4. 4	. a	1.78	1.27	1.02	•
850	743452	AA609368	0.58	7 .	2.64	1.47	0.50	0.0	2.87	1.46	1.20	•
921	289706	N62948	0.33	0.87	2.63	2.06	6.78	- t- c		1.71	8.55	-
922	505538	AA147594	0.09	0.25	2.63	2.25	0.11	2:32		. 69	2.86	-
923	53022	R15880	0.89	2.34	2.62	2.59	1.25	1.32	5.6	9 c	19	-
924	273546	N33274	2.51	6.55	2.61	2.25	2.58	2.24	6.05 4.05	0.60	1.45	
20	843049	AA485983	0.10	0.25	2.61	2.59	1.11	8:20	1.27	7.7	7 7	
300	504517	AA189845	31.25	81.32	2.60	1.36	2.19	2.32	6.51	9/.	o	- •
276	45347K	AA700218	66.0	2.58	2.60	5.15	2.24	1.63	2.78	1.77	2.03	- •
3 8	20000	A A 042255	0.28	0.67	2.60	1.94	9.92	1.20	1.46	0.37	0 6	- ,
9 6	100000	AA01333	3 6	10.21	2.58	2.40	2.96	1.23	5.65	1.74	56.1	- •
878	070040	15421W	3 5	033	2.58	0.08	0.69	12.77	0.53	0.28	1.15	- ,
3 3	4/7440	W/3011	2 6	3 6	2.58	9.39	0.82	1.72	2.27	0.98	0.31	- ,
3 8	7CI RRC	60155744	3 5	2,65	2.58	1.48	1.68	2.03	2.47	5.56	2.26	- ,
723	045/0/	AA416504	3 6	3 6	2.58	0.98	2.46	5.63	1.51	1.91	2.98	- ,
933	743727	AA629355	0 0	77.0	2.57	1.64	1.73	1.89	1.42	7.19	1.57	-
8	810625	AA464755	0.07	5 5	7. C	198	1.32	1.27	7.42	2.20	1.24	-
935	1476181	AA872034	0.20			- 5	1.20	1.67	8.80	0.59	1.96	-
936	814154	AA496252	0.87	2.23	2.5/	1.20	} -	1.07	0.94	0.72	10.42	₩-
937	34832	R45157	0.59	1.53	2.57	5	14	264	1.01	0.54	7.34	-
838	435820	AA701519	0.50	1.29	2.56	2.29	/o:1	1.35	2.96	1.57	2.33	-
838	130703	R22003	3.70	9.47	2.56	08.1	23.5	920	2.28	2.59	5.66	•
g S	30428	R42061	3.15	8.00	40.2	67:1		2 98	88.8	0.36	1.25	-
ጀ	299162	N75473	0.03	0.07	2:54	0.36	0.30	7 7	2.09	1.48	2.38	-
3	267679	N25552	0.58	1.47	2.53	2.42	8°.0		88 7	8 28	0.17	•
253	145383	R77919	0.10	0.24	2.52	1.76	1.91	2. ·	3 6	186	1.60	-
3	154720	R55220	0.05	0.13	2.52	2.29	7.29	1.91	BL:0	5.00	7	
2	395955	AA757564	0.49	1.22	2.52	1.20	1.50	1.33	0.70	- 6	+ - C	
8	412949	AA707840	0.92	2.31	2.51	1.63	5.07	2:31	1.74	80.	3 6	٠,
3	120892	Tosaga	3.38	8.48	2.51	1.12	1.30	2.13	,	1.86	0.02	- •
8	271989	N31935	0.18	0.45	2.50	1.45	2.91	1.52	1.12	5.41	76.3	- •
3	692592	AA159577	3.36	8.37	2.49	1.59	0.54	2.12	9.45	0.77	4.	•
}			:				28					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold of higher in at least one fund sample or 3 fold or higher in at least 2 tumor samples.

		2 2 2 2			:		COLUMN TO MANAGEMENT OF TA	Fold/MPM124/A	Fold(MPM122/A	Saldine Mandalla Foldine M124/A Foldine M122/A Foldine M123/A Foldine M123/A Foldine Foldine	old(MPM125/A	# of Tumor
	IMAGE	Accession	AveBPH	AveTumor	Fold(AveTumor/Av eBPH)	Fold(MPM12U/A	veBPH)	veBPH)	veBPH)	VeBPH)	veBPH)	Samples
		Number			•	ě	70.	2.4.0	1.57	1.07	1.62	-
920	591055	AA161097	0.08	0.20	2.49	7.00	¥7:-		1.50	1.97	1.31	-
951	267252	N24579	0.83	2.07	2.48	6.02	2.69	9°.		121	9.12	-
52	53393	R16245	1.1	2.75	2.48	1.34	1.22	F8:0	9	. 88	0.48	-
	327058	AA284267	0.16	0.39	2.47	9.38	1.43	0.13	 	3 6	1.48	-
3 3	204544	NG0425	0.51	1.25	2.47	2.70	5.07	2.03	2.61	t 9	2 6	•
	70000	7007450	28.85	70.79	2.47	2.13	1.92	1.40	5.51	08.7		
	£08707	104050	200	0.87	2.47	1.93	2.00	1.57	4.15	5.60	r (1)	- •
926	112456	T80871	77.0	6.0	2.47	1.81	1,94	1.23	1.48	6.78	1.58	- ,
957	810779	AA481754	0.20		2.6	£ 5	5.16	2.27	1.48	2.27	1.99	- ,
958	431933	AA678164		2.73	04:7		2.24	2.13	1.46	1.27	6.81	-
959	359653	AA010872	0.32	0.80	2.46	0.0		1.59	5.29	1.80	2.03	-
8	681865	AA256157	0.61	1.51	2.48	1.43	10.7	90 0	1.06	0.89	1.07	-
961	239199	H70942	0.91	2.25	2.46	0.90	67.0		78.0	1.05	1.99	-
962	66564	T67058	0.31	0.76	2.48	2.60	7.66	0.0	. a	2.82	5.77	-
583	215000	H73241	3.61	8. 48.	2.45	2.77	1.25	0.82	2 ;	4 5 F	1.22	-
3	840773	AA481751	0.87	2.12	2.44	1.12	1.19	8.67	/L.F	2 .		_
\$	2000	1011011	6	1 20	2.44	2.00	1.90	5.73	1.70	79.	2 C	٠.
9	78700	OFFICE C	3	. A	2.42	1.35	2.68	2.96	5.20	1.37	0.80	- •
3	201301	K885/3	, i	2 9	2.42	1.55	1.08	2.	0.90	9.27	99.0	- ,
267	773170	AA428514	0.17	2 i	7 (7	2 88	6.50	0.47	1.05	1.75	2.16	-
88	897656	AA496796	5.26	12.74	74.7	8 4	1.71	2.00	1.45	2.38	5.02	•-
68	111884	T84996	0.03	0.02	2.41	80 1		2.43	5.19	2.25	1.49	-
970	744897	AA625784	1.69	₹	2.40	/6.1	P 6	78.0	5.14	1.19	1.88	-
178	46411	H09164	3.30	7.91	2.40	1.72	1.70	5 6	2 13	2.35	1.52	-
972	796761	AA460729	0.59	1.42	2.40	1.08	رة. ا	0.00	<u>.</u>	08.0	1.70	-
973	1031273	AA609054	0.38	0.91	2.39	1.15	7.94	80.L		08.90	2.65	-
974	46129	H09529	0.07	0.18	2.39	0.81	1.22	1.80	48 48	7.40	2.27	-
975	220389	H86816	1.1	2.65	2.39	1.00	1.09	1.16	e 4	25.	0.88	-
978	1292170	AA705819	1.48	3.50	2.38	1.36	1.60	8.89 18.00 1	00.1		1.68	-
277	356711	W84580	0.78	1.86	2.38	1.32	2.39	1.02	1.38	. v	4	-
	120884	A 6.78203	0.65	1.56	2.38	1.54	2.21	2.38	9.00 40.00	07:1	<u>.</u>	٠ •
	25575	DAFORA	2.28	5,43	2.38	1.71	0.91	2.32	2.25	1.78	97.6	- •
		1000		1.26	2.37	0.80	1.53	1.02	8.43	-	5 0	- ,
	2000	80171		2 2	2.37	0.89	0.90	0.69	0.41	10.29	c0.r	- ,
200	CR59951	AAS44141		* *	2.37	1.39	1.02	8.38	2.45	0.70	0.28	- ,
28 8	58//81	K83/58	0.70 7.85		2.37	0.64	1.43	1.12	99.9	2.63	1.71	-
Š	20130	RORSY	3	: •			29					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one funder sample or 3 fold or higher in at least 2 tumor samples.

	٠									A/204118111/17-1	TOIM MEM 125/A	* of Tumor
	IMAGE	Gen Bank Accession	AveBPH	AveTumor	Fold(AveTumor/Av Fold(MPM120/A Fold(MPM124/A	old(MPM120/A l	Fold(MPM121/A F veBPH)	old(MPM124/A veBPH)	-old(MPM122/A veBPH)	veBPH)	veBPH)	Samples
		Number	;	;	900	. 96	0.96	1.27	8.99	1.09	0.62	- -
8	39899	R52624	0.60	1.41	0.30	7.56	0.57	2.63	1.43	1.70	0:30	-
985	276815	N34824	0.13	0.32	00.30	2 6	2.92	0.89	0.84	0.87	1.75	-
988	430007	AA034115	0.48	- -	2.35	6.0	1.59	1.65	1.23	1.13	1.21	-
287	121615	T97592	0.4	. 9.	2.35	1.5.		1.13	1.43	7.48	1.30	-
888	826303	AA521016	0.54	1.28	2.35	67.1	S 6	2.43	8.64	0.85	0.41	-
888	3442	R44985	1.70	4 .00	2.35	99.0	5 6	1.52	41.1	5.44	2.13	-
068	742542	AA400010	0.20	0.46	2.34	1.7	2.12	 	533	1.78	2.68	-
99	327678	W23758	0.30	0.70	2.34	1.23		2 6	1.18	1.22	9.40	-
885	160658	H24954	0.07	0.17	2.34	0.52	4.0	. q	5	9.44	2.14	-
883	265114	N20820	0.34	0.80	2.34	7.86	68.0	0 0	- 62	980	6.28	-
8	626842	AA191424	1.69	3.95	2.34	1.72	1.97	7:27	6.0	2 12	2.12	
588	291403	N87762	9.0	0.10	2.33	5.38	0.72	2.23	\$ i	7 7 7		. 🕶
8	240318	H89795	0.43	9.	2.33	1.83	2.67	<u>.</u> Ž	9.0°	0.0	87 7	•
8 6	252884	H88334	4	3.82	2.33	0.77	0.58	9.21	0.93	88.0	<u>.</u>	- 🕶
9 0	302400	WHR724	0.18	0.41	2.33	1.08	1.73	2.18	1.05	6.70	5.00	- +
	420742	520024	283	9	2.33	1.55	1.60	7.25	1.31	1.29	0.80	- ,
		A 820011	, ,	305	2.33	0.67	0.64	0.61	9.75	1.41	0.87	
3 3		AA6624811		9 6	2.32	1,41	1.54	1.91	1.	1 .54	6.13	- ,
3		AA002168	2 0	3 6	232	2.51	5.63	1.48	1.28	1.99	1.05	 ·
1002		H98244	6.0		333	5.12	1.92	2.46	2.71	0.86	0.85	-
<u>8</u>		AA758470	90.0	2 ;	7.77	1 41	0.59	0.79	9.48	0.33	1 .34	-
ş		AA001538	0.46	1.07	2.32	- 60 - 6	1 17	8.26	1.19	1.30	1.05	-
1005		AA192765	2.16	86.4	2:32	2 20	0.79	1.98	2.04	2.01	5.13	-
8		R42831	0.03	90.0	2.30	t 90:-	2.6	0.17	2.06	9.31	0.70	-
1001	,	AA005228	0.25	0.58	87.7	0.90	1.43	7.27	1.22	1.36	0.83	-
8	•	AA608952	t 5	S	9 00 0	30.0	. 38	1.79	6.04	2.82	0.65	
60		AA427927	18.90	5.54 5.55 6.53	87.7	5 6 6	1.72	<u>5</u>	5.27	2.14	2.50	- -
5	••	N90051	0.92	01.2	6.40	3 6	2.64	2.00	1.62	1.22	1.18	-
194		R41685	r	0.28	07:7		, og	1.89	1.85	6.37	1.85	-
1012		AA598520	0.68	C .	2.27	0.80	1.27	1.75	2.28	5.09	2.57	-
193	430052	AA034179	0.40	78.0	77.7	. i	77	2.71	1.32	0.42	2.18	-
101		H10665	0.17	0.38	2.27	, s	17.0	0.71	1.69	0.62	9.05	~ -
1015		AA278749	8.47	19.24	2.27	3 6		0.87	1.58	9.54	1.19	-
1016	••	W68396	0.42	0.98	2.27	2.0	18.5	2.33	2.50	0.73	1.15	-
1017	25156	R38935	1.15	2.61	77.7	<u>.</u>) 				
							30					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one fumor sample or 3 fold or higher in at least 2 tumor samples.

	IMAGE		0	YounTenA	Fold(AveTumor/Av Fold(MPM120/A Fold(MPM121/A Fold(MPM122/A Fold(MPM123/A Fold(MPM125/A # of Tumor	Fold(MPM120/A	Fold(MPM121/A	Fold(MPM124/A	Fold(MPM122/A veBPH)	Fold(MPM123/A I	Fold(MPM125/A veBPH)	# of Tumor Samples
	Clone ID	Number			•BPH)	Vebru)			2	0.47	6.54	-
9	730734	4444807	0.08	0.18	2.26	1. 4	2.38		2 6	630	2.58	-
5 5	tooto,	A A DEFOTO	800	0.17	2.25	2.09	0.54	1.89	5.0	7 7	0.17	•
BLOL	1/0//5	AA055878		, A	2.25	0.33	0.35	0.39	61.11	7 7	175	•
1020	271799	N31605	87.7	2 3	2.25	0.83	1.65	0.93	6.21	11.2	5 6	٠ ٦
1021	448885	AA777670	5.05	٥٠. د		7	0.94	2.83	5.99	1.42	R O O	- ,
1022	171836	H19068	0.37	0. 8 . 4	2.24	5. 6	1.43	96.0	1.26	8.84	4.0	, .
1023	379709	AA778089	1.21	2.70	2.23	0.40 0.40		6.10	2.62	1.52	1.65	-
1024	325012	AA284283	0.09	0.21	2.23	0.18 8	87.1 60.0	7 00	1.27	1.25	1.20	~
1025	363058	AA019316	0.22	0.48	2.22	7.67	0.92	5 6	1.35	1.81	1.21	-
1028	48914	H10208	0.78	1.73	2.22	6.51	14.	8 8	2	1.07	2.97	-
1027	23819	R39446	0.38	0.85	2.21	0.63	1.56	1.02	73 7	1.36	1.27	-
10.2	772960	AA476257	0.54	1.18	2.21	1.60	2.11	0 0		10.80	0.55	-
1028	71730	T51229	0.0	0.20	2.21	0.08	0.07	0.0	8 6	20.	2.16	-
5 5	281571	H98756	0.03	0.07	2.20	1.59	5.46	0.08 0.08		0.27	0.27	-
2 5	814053	AA465495	10.76	23.71	2.20	99.0	0.35	2.52	9 6	7 63	1.13	-
2 5	4202887	A A 778755	0.48	1.05	2.20	1.24	1.46	1.02	0.71	80:-	1.59	-
7007	1007671	A 2284278	0.25	0.54	2.20	1.76	1.64	5.61	09.1	2.60	1.30	-
3	324000	TERROR	2	8	2.20	0.86	1.78	1.92	5.87	<u>}</u>	A 24	-
1 03	72000			, to t	2.19	1.91	2.16	1.40	0.97	1.53	- 84	. 🕶
1035	1467109	`	70.0	07:		0.59	1.13	1.73	0.09	0.0 40.0	0 00	• •
1038	244951	N54551	0.25	0.55	2. t	22.0	0.65	9.52	0.83	0.76	0.62	- ,
1037	593280	AA165512	98.	4.07	81.2 61.5		1.66	1.99	1.69	1.34	5.45	-
1038	296040	N73584	1.50	3.29	2.19	1.03	3 5	96	0.50	0.22	10.37	-
1039	451871	AA706935	0.16	0.35	2.18	0.69		6.35	141	1.27	1.67	-
5	795588	AA459692	0.78	1.70	2.18	1.13	07.1	0 to	6.01	1.14	1.76	-
140	166199	R88246	0.05	0.11	2.18	1.50	0.53) w	0.12	0.12	2.07	-
1042	46183	H09616	90.0	0.18	2.18	1.76	74.0	70.7	1 70	1.05	6.35	-
1043	344806	W72838	0.42	0.91	2.18	1.28	1.47	t 90 0	701	0.55	0.73	-
124	51939	H22958	0.42	0.91	2.18	8.97	0.79	0.90	5 5	98.0	0.19	-
1048	28082	N50808	0.05	0.11	2.17	10.72	0.90	0.18 8.1	<u>a</u> 6	8 4v	1.19	-
2 5	4404070	A \$ 800447	0.55	1.09	2.17	1.57	1.72	1.25	0.82	? tr	1.1	-
	0/010#1	7408014V	5 6	6	2.17	2.12	2.37	0.25	0.25	0.70		•
3 5	400000	188343	5 6	1.52	2.16	1.47	1.12	1.37	1.17	ο ο	, v	
\$	086717	315000	7	273	2.18	2.42	1.28	0.74	0.85	5.42	17.7	• •
1049	147834	R81831	2 6	2 6	i c	0.42	0.39	1.55	0.49	0.57	LC:30	- ,
1050	418155	W86075	0.23	8 t	5 t	1.94	7.04	1.24	99.0	0.70	1.32	-
1051	139837	R62289	40.0	<u></u>	ì		7					

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		725				ALCO PREMIENTS OF THE	COLUMN TANDERS OF A CA	Fold(MPM124/A	old(MPM122/A	Fold(MPM123/A F	old(MPM125/A	# of Tumor
	IMAGE	Accession	AveBPH	AveTumor	Fold(AveTumor/Av eBPH)	FOIG(MFM120/A)	veBPH)	veBPH)	veBPH)	veBPH)	veBPH)	Samples
		Number			. !	69 6	1 32	1.95	1.79	6.30	0.92	-
1052	131626	R23735	0.10	0.22	2.15	0.00	3 5	4. t	7.30	1.06	0.88	-
1053	32663	R43319	0.50	1.07	2.15	0.89	74.1	÷ •	x 23	1.14	1.10	•
1054	1031717	AA609584	1.03	2.22	2.15	1.83	6 <i>)</i> :	9 6	27.0	0.79	0.60	-
1055	254328	N22272	1.39	2.97	2.15	0.66	0.43	o 0		0.52	1.00	-
1058	788605	AA452897	99.0	1.42	2.14	1.48	2.24	2.10		9	2.64	-
1057	166245	R88267	0.03	0.07	2.14	0.31	5.65	15.0	10.3	0,00	0.38	-
1058	32991	R44770	0.26	0.55	2.14	10.49	0.04	1.10	0.6	8 1 1 1	1.08	-
1059	712840	AA282023	0.13	0.28	2.13	1.31	0.35	.	æ !		200	. 🕶
1060	359153	AA010136	0.47	9.5	2.13	1.35	1.22	6.15	1.71	\chi.	0.92 7.87	
8	207864	H50581	0.42	0.89	2.13	1.08	1.48	2.05	1.23	<u>\$</u>		٠ +
5 5	305711	AA757764	0.10	0.22	2.12	1.66	5.61	1.05	1.68	35.1	0 7	- +
§	78948	TR1792	3.51	7.43	2.12	1.23	0.53	1.70	1.68	5.T	- ¢.	- 🕶
\$ \$	42373	B67147	2.61	5.52	2.11	2.11	0.91	1.90	1.25	 		
8	199628	20850R	0.18	0.37	2.10	1.07	1.40	0.88	1.69	0.70	9 c	- •
8	29828	R42331	1.30	2.72	2.10	2.03	0.72	0.48	2.13	3 . 6	5.32	- +
5 5	809828	A4455571	0.66	1.39	2.10	6.79	1.73	0.53	1.28	68:0		- •
3 8	2000	NESSTR	08.0	88	2.10	1.31	1.57	0.58	1.09	0.95	01.7	- ·-
3 8	78408B	0 4 4 3 0 0 0 3	90	131	2.10	1.23	0.99	6.61	1.56	1.28	0.92	- •
8 9	000101	114091	5 6	5	2.09	1.05	0.99	1.90	1.23	1.17	6.21	- •
	70002	19BICN	5 6	3 3	2.08	0.47	1.67	0.47	1.20	7.98	0.69	. .
	6/079/	AA44/585	7 7	, ,	202	1.21	1.21	1.49	5.44	1.88	1.22	_
1072	73785	154672	S	8 3	20.5	2.83	1.29	0.05	2.06	5.24	0.73	-
1073	730124	AA412470	77.0		 	38	0.99	0.32	1.09	7.73	0.79	-
1074	51483	H18924	07.0) i		. 790	1.05	0.28	0.36	9.67	0.30	-
1075	38487	R49124	\$? • •	27.5	2.03	1.67	1.45	1.18	1.05	6.22	0.63	-
	27000	AM020042		t 4	2 03	0.45	0.67	0.71	0.58	8.86	0.91	, .
701	010141	K/0003	3 6) E	090	0.98	0.58	2.60	1.86	5.54	-
9/01	15/33/	AA437089	9 5	9 6	. c	7.29	1.20	0.95	0.79	1.68	0.22	-
8/01	ACOCC7	N2/041	3 6	3		0.05	0.31	0.75	0.65	89.6	0.69	-
2	428169	AA001876	0.20	.	2.02	8 6	118	6.48	1.57	0.85	0.92	-
108	950470	AA599102	2.48	9.00	2.02	5 4		0.92	1.59	6.92	1.39	
1082	824044	AA491206	0.09	0.18	2.0,1	- 6 - 6		130	5.81	1.46	0.49	- -
1083	1458721	AA864875	12.97	26.09	2.01	1.29	707	58.0	0.37	8.19	0.71	-
<u>\$</u>	757158	AA443938	0.49	0.98	2.01	0.80	17.1	3 6	80	0.87	0.98	-
1085	252830	H88261	1.63	3.26	2.00	1.12	0.93	47.1		; ;		
							cc					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

	HAGH	Gen Bank	į		Fold(AveTumor/Av	Fold(MPM120/A	Fold(MPM121/A	Fold(MPM124/A	Fold(MPM122/A	veTumor/Av Fold(MPM120/A Fold(MPM121/A Fold(MPM124/A Fold(MPM122/A Fold(MPM123/A Fold(MPM125/A ≢ of Tumor/Av Fold(MPM120/A Fold(MPM121/A Fold(MPM124/A Fold(MPM122/A Fold(MPM123/A Fold(MPM125/A ≢ of Tumor/Av Fold(MPM120/A Fold(MPM121/A Fold(MPM124/A Fold(MPM122/A Fold(MPM123/A Fold(MPM125/A ≢ of Tumor/Av Fold(MPM120/A Fold(MPM121/A Fold(MPM124/A Fold(MPM122/A Fold(MPM123/A Fold(MPM125/A ≢ of Tumor/Av Fold(MPM120/A Fold(MPM121/A Fold(MPM122/A Fold(MPM123/A Fold(MPM123/A Fold(MPM125/A ≢ of Tumor/Av Fold(MPM120/A Fold	old(MPM125/A	# of Tumor
	Clone ID	Accession	АувВРН	AveBPH Averumor	eBPH)	veBPH)	veBPH)	VeBPH)	(mag)	:		•
		Lagun.	#		50 6	5.45	1.50	2.33	0.65	0.59	1.50	-
1086	24918	R39069	2.08	o :	0.7	· ·	001	1.62	5.66	1.33	1.27	~ ~
1087	1637296	A1005519	25.34	50.73	2.00	2 9	50.	6.39	1.19	1.19	0.99	
1088	123539	R00809	0.73	- .	2.00	0.1	5 6	22.0	1.57	5.84	0.29	
1089	1461664	AA885311	2.34	4.66	1.99	1.88	6. 4	 	. 6 	1.30	0.58	· -
1090	1467207	AA884713	5.60	1.14	1.99	1.05	10:1	99°	1.76	0.57	6.35	-
1091	79022	T62179	7.89	15.78	1.98	2.00	0.40	1.28	1.57	0:30	6.37	•
1092	37665	R61374	0.03	90.0	1.96	1.33	0.87	1.50	5. 50	0.52	1.47	₩-
1093	452545	AA778839	0.57	1.12	1.96	1.27	1.70	1.07	0.83	7.14	0.09	-
1 08	773568	AA428196	0.12	0.23	1.95	0.56	6.03	79.0	1.65	7.29	0.49	-
1095	480894	AA136699	0.35	69.0	1.95	0.70	6.6	9.53	5.73	0.84	2.42	-
1096	41074	R56604	0.21	0.40	1.85	0.77	04.1		, ,	1.50	0.86	-
1097	257342	N29638	0.51	0.89	1.95	1.38	1.30 0.50	0. 4 0. 4	0.00	1.63	5.75	-
1098	897595	AA496887	0.85	1.66	1 .	1.40	0.89	0.10	6.00	0.57	6.65	
1099	108378	T77729	0.02	0.03	2 .	2.53	0.85	0.47	9000	8.19	1.03	*
5	276911	N34945	0.21	0.40	<u>2</u> .	0.63	0.65	0.73	0.50 0.00	2.74	0.13	-
101	147075	R80235	0.07	0.14	1.93	0.12	5.30	 	0 t. 4	1.27	0.71	-
1102	773331	AA425436	10.48	20.23	1.93	0.68	0.73	2.3	5.00 8.4	1 48	0.69	-
1103	241077	H80325	3.25	6.25	1.93	0.86	1.38	o	0.0	333	0.57	-
<u>‡</u>	824074	AA491227	0.18	0.3 4	1.92	8.92	0.68	6.52	 	0.11	6.21	-
1105	796078	AA480353	0.09	0.18	1.92	1.61	0.75	1.07	1.20	1.12	0.85	-
1108	838616	AA456968	1.21	2.31	1.91	0.59	0.58	9 4 ./) 8	1 81	0.48	•
1107	45501	H08397	0.12	0.22	1.91	2.63	2.45	0.03	99.6	0.00	2.40	.
1108	415279	W95050	0.51	0.97	1.90	1.12	0.89	40.0	08.0 84	38.0	0.83	-
109	40299	R52085	0.10	0.19	1.89	1.50	2.20	0.80 0.00	6.5	7.10	0.33	-
110	33894	R44530	0.26	0.48	1.89	0.99	0.03	7 8	9 7	08.0	0.95	* -
1111	471641	AA034945	4.72	8.92	1.89	90.9	1.75	0.60		4.6	0.72	-
1112	742887	AA406226	0.36	0.68	1.89	1.23	87.1	97.7		150	0.60	. -
1113	346523	W73994	1.0	1.90	1.89	2.65	5.94	1.07	0.92 7.87	1.57	1.79	,
1114	743229	AA400329	0.36	0.67	1.88	0.82	0.76	5 to 1	5 6	75	0.25	-
1115	866702	AA679180	0.0 4	0.08	1.88	0.25	5.80	1.45	88 (27.0	1.29	-
1118	150798	H02550	1.20	2.25	1.87	0.95	6.61	88.0	7 7	0 C	102	, -
1117	183556	H44032	0.28	0.52	1.87	5.21	-	5 .	? •. •	- u	80.0	
1118	243905	N33965	0.19	0.35	1.86	0.27	1.33	2.60		9 6	27.0	
1119	731404	AA412247	0.92	1.72	1.86	1.31	0.90	5.72	1.23	77.) ;	•
							33					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumorisample or 3 fold or higher in at least 2 tumor samples.

		4			September 12 Total Control Con	A100	ECIMINATION A	Fold/MPM124/A	Fold(MPM122/A	Fold(MPM123/A	Fold(MPM125/A	# of Tumor
	IMAGE	Accession	AveBPH	AveBPH AveTumor	Fold(AveTumor/A)	/ Fold(mFm120/A VeBPH)	veBPH)	VeBPH)	veBPH)	veBPH)	VeBPH)	Seldings
	Cione	Number					80 0	1.02	5.71	1. 29.	0.69	-
1120	243321	H94720	1.08	2.00	1.86	0.92	0.90		 	1.05	5.57	-
121	243803	N49725	9.0	1.19	1.86	1.16	1.11			1.67	0.17	•
1122	841610	AA487457	90.0	0.11	1.86	1.70	5.30	2.13	3 6	121	1.09	-
<u> </u>	80374	T85765	1.21	2.24	1.85	0.50	6.17	T.4.	9 9		0.67	-
12,4	731029	AA421269	0.08	0.15	1.85	1.68	2.25	0.69	9.70	2.61	0.92	-
1125	284882	N66737	0.33	0.61	1,85	0.44	0.82	0.56	t (i	57.0	0.26	-
, t	1031009	AA635186	5.84 4	10.78	1.85	0.60	0.46	1.52); ;	7	1.13	•
1427	1030637	AA608770	0.76	1.4	1.85	1.20	1.32	5.23		. r	0.0	•
4128	273168	N36794	0.15	0.27	<u>48.1</u>	1.65	0.62	9.64	2.12	900	1.28	. 🕶
130	68317	T8816	8.85	16.25	1.84	6.02	2.40	0.49	7 7 7	08.0	0.92	-
	253934	N22084	1.1	2.03	1.83	0.86	0.96	6.40	5 3	9 6	1.05	-
3 5	48708B	A A DA 5300	2.58	4.69	1.82	0.88	0.91	5.53	* 6		1.68	-
<u> </u>	267676	0000000	1.38	2.50	1.81	0.86	0.87	5.81	0.92		0.61	
3 5	4068488	A 8 2 1 0 0 1	0.86	75	1.79	96.0	1.47	5.27	1.26	0.50	0.80	-
3	0010001	4 4 00 6 0 0	34 DK	57.14	1.79	1.13	0.53	1.83	5.43	0A:0	1 2	•
<u>*</u>	160683/	AASSOUZO	G 10	1	1.78	0.21	5.41	1.10	2.06	0.17	7.7.	- +
133	343731	W69213	9 6	- 4	178	0.12	0.55	1.10	7.92	0.89	21.0	- •
138	25058	R38894	0.0	<u>.</u>	7 -	68.0	1.40	1.21	5.01	1.13	1.03	- ,
1137	257387	N30704	2.69	/s		08.0	0.64	6.31	0.75	1.01	1.13	- •
1138	128058	R09729	1.13	2.00		28.0	76.0	0.61	0.83	6.53	0.83	-
1139	397604	AA701034	0.18	0.32	1.76	9	0.50	0.88	0.50	0.50	6.13	-
140	42271	R60328	0.05	0.0 2	1.76	5.00 5.00	1 6	1.28	0.30	0.30	1.28	-
##	71674	T57920	0.24	0.42	1.76	9.0 40.0	ţ ;	7.79	2.19	2.5	90.0	-
1142	340949	W57818	0.17	0:30	1.74	0,46	- G. W	0.0	0.10	2.82	0.10	-
143	788256	AA454098	0.10	0.17	1.73	0.19	02.6	80:-	78.0	0.50	5.93	-
+	142397	R69934	99.0	1.13	1.72	0.90	00:L	9:1	35.0	1.28	1.26	•-
1145	66902	T69477	0.89	1.53	1.72	0.49	6.39		503	1.66	0.71	-
1146	26506	R20640	0.47	0.80	1.72	96:0	C/:0		000	6.93	1.23	-
1147	489476	AA045673	0.11	0.20	1.72	0.86	60:0	0.31	6.00	0.09	0.00	-
1148	795446	AA453616	0.12	0.20	1.72	1.17	2.30	9.68 70		0.85	0.51	-
1149	471498	AA035347	90.0	0.14	1.71	0.44	5.02	1.2.1	4.2.3 67.3	0.93 0.83	0.35	-
1150	745001	AA626011	1.43	2.45	1.71	0.67	1.30	1.30	3.72	80.8	0.78	-
1151	609155	AA176867	2.10	3.59	1.71	0.49	1.44	0.00	9 6	0.72	0.72	•
1152	796549	AA460274	0.01	0.02	1.70	0.72	0.54	9.00	9 6	5.68	1.23	-
1153	363003	AA018866	0.40	0.68	1.70	1.05	1.03	0.60	0.00	3		
							34					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

						Fold(MPM122/A Fold(MPM123/A Fold(MPM123/A Fold(MPM123/A # of Tumor	V/707114111111111111111111111111111111111	E.I.A.MDM124/A	Fold/MPM122/A	Fold(MPM123/A	Fold(MPM125/A	# of Tumor
	IMAGE	Gen Bank Accession	AveBPH	AveTumor	Fold(AveTumor/Av	, Fold(MPM120/A	Fold(MPM121/A	veBPH)	VeBPH)	veBPH)	veBPH)	Samples
	Clone 10	Number				•	,	4 0 0	0.83	6.01	0.14	-
7377	974183	N21808	0.07	0.12	1.69	1.75	0.14	1.40) ; ;	280	0.78	-
5	20117	000001	2 62	4.28	1.69	0.88	0.60	1.14	00.0	9 6	71.7	-
1155	46350	000000	3 4		1.69	1.60	0.45	0.84	1,35	2.0		•
158	825461	AA504354	S :	8:-	9 7	171	1.08	0.81	0.41	re.o	<u> </u>	• •
1157	1291950	AA707450	0.32	0.55	80.1		990	6.32	0.82	0.78	0.50	
1158	418197	W90364	96.0	1.60	99:) (i	0.00	1.20	5.19	9.0	0.62	-
1159	133358	R26859	0.61	1.01	1.66	7.52		0.38	<u>¥</u> .	5.30	0.18	-
8	269606	N26769	0.06	0.10	1.65	0.27	. t. t	102	0.23	99.0	5.01	- -
1161	1293119	AA682233	0.46	0.76	1.65	1.60	. S. C.	98.0	1.09	7.12	0.50	-
1162	768953	AA424920	2.45	4 .03	1.65	0.50	0.28	0.73	0.23	0.16	7.41	-
1163	772916	AA479912	90.0	0.10	1.	0.16	0.10	0.10	0.74	5.35	0.35	-
1164	31869	R43017	0.23	0.37	1.62	2.56	0.27	2 6	72	0.97	9:3	-
1165	84141	171042	0.03	0.05	1.62	2.59	5.16	4, 64 64, 64	50.00	2.20	0.97	-
1168	969636	AA663592	1.87	3.03	1.62	0.64	0.38	0.00	7 87	0.18	0.34	-
1167	810697	AA457696	0.14	0.22	1.61	0.07	0.07	0.10) () ()	0.14	0.68	-
1488	37234	R35283	0.07	0.11	1.61	1.33	1.88	0.0		0.11	2.13	-
3 5	186132	H39580	0.24	0.38	1.59	1.04	0.41	0.24	20:0	46.	0.10	-
	4428R	T01168	0.10	0.16	1.59	1.13	0.87	1.25	0.02	69 4	0.09	-
2 ;	100447	A A 48.0081		0.18	1.58	0.29	1.78	0.70	0.80	9 6	5.73	-
1711			. 6	41.0	1.57	1.21	1.33	0.95	0.11	- i		-
1172	41321	CORROCA	8 6	6. 6.	1.57	0.26	0.65	0.32	1.31	0.75		
1173	429446	AA007819	0.50	7 9	155	0.89	5.96	1.61	0.09	80.0°	3 3	· •
1174	882571	AA676515,	E :	- (- - (-) ¥	1.72	0.40	0.73	0.61	0.40	5,41	- •
1175	448813	AA777590	1.22	1.89	20.	35	1.37	0.76	6.22	0.22	0.33	- ,
1176	49358	H15104	0.22	0.3 %	đ. (990	0.36	0.46	0.86	6.51	0.31	.
1117	810891	AA459519	0.36	0.55	7.52	0.00	0.75	5.42	0.73	0.70	0.68	- ,
1178	594123	AA169475	1.91	2.89		5 6	7.25	0.59	0.31	0.31	0.31	-
1179	869375	AA679907	0.03	0.05	1.51	. c. d	65.0	0.11	0.88	0.62	0.27	-
1 180	128248	R11510	0.09	0.14	1.50	6.9	20.0	0.44	0.51	5.78	0.55	-
1181	22895	R38640	0.09	0.14	1.50	i !	500	5.33	0.81	0.65	0.71	-
1182	286404	N67274	1.50	2.22	1.48	0.77	7 00	57.0	134	5.61	0.31	-
1183	366915	AA027147	0.16	0.23	4.	0.60	90.0		0.20	0.25	0.25	-
1184	591465	AA160695	9.0	90.0	1.42	0.40	\$0.0 \$0.0	5.5	80	44	0.63	•
1185	344595	W73634	0.37	0.52	1.41	5.41	0.92	t 70 0	0.31	78.0	6.29	•-
1188	781110	AA430042	0.03	9.0	1.38	0.23	0.31	5.0	523	0.24	5.97	-
1187	291690	N67832	0.0	90.0	1.38	0.16	0.20	98.0	5			
!							35					

Samples 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
Fold(MPM126// veBPH) 0.49 0.07 5.71 0.46 0.91 0.45 0.45 6.44 0.33 5.28
Fold(MPM123/A veBPH) 5.54 0.07 0.11 0.27 0.18 5.08 0.29 0.47 0.12
veBPH) veBPH) 0.42 0.56 0.75 0.75 0.76 0.76 0.41 5.73 0.43 0.67
old(MPM124/A 1 veBPH) 0.53 0.26 0.14 0.18 0.26 0.30 0.44 0.14
Fold(MPM121/A F veBPH) 0.52 0.76 0.46 0.29 0.44 0.45 0.12 0.12
Fold(MPM120/A veBPH) 0.74 6.50 0.60 0.62 0.43 0.45 0.12 0.12
Fold(MPM12xIA VeBPH) VeBPH) Post NeBPH VeBPH V
AveTumor 1.44 0.20 0.11 2.37 0.09 0.07 1.83 3.64 0.10 0.10
AveBPH 1.05 0.14 0.08 1.78 0.07 0.06 1.41 2.86 0.17 0.09 0.16
Gen Bank Accession Number AA670390 AA875953 AA129677 AA048430 R60722 H29216 AA688959 N95226 AA688959 N95226 AA688959 N95226 AA60670 W333343
IMAGE Clone ID 878469 1482258 565493 42118 52594 854401 307337 377296 592728 322617
1188 1190 1190 1192 1193 1194 1196 1196

Table 2-2: T3NO vs. BPH: Gene upregulated 5 fold to higher in at least 2 tumor samples or 3 fold or higher in at least 3 tumor Samples.

Table 2-2: T3NO vs BPH: Genes upregulated 5 fold or higher in at least 2 tumor samples.

Accession Designation Availability Foliative contains Availability Foliative contains Availability Foliative contains Availability Availability <th< th=""><th></th><th></th><th></th><th></th><th>186</th><th></th><th></th><th></th><th>-</th><th>A/401011011494/A</th><th>Entering M124/A</th><th>Fold/MPM122/A</th><th>Fold(MPM123/A</th><th>Fold(MPM125/A</th><th># of Tumors</th></th<>					186				-	A/401011011494/A	Entering M124/A	Fold/MPM122/A	Fold(MPM123/A	Fold(MPM125/A	# of Tumors
788180 AAASS310 proteins	Je pe	IMAGE	-	Tissue	mbrane	AveBPH	AveTumor	Fold(AveTumor/A	rold(MrM120/A	veBPH)	veBPH)	veBPh)	∨ •BPH)	Ve@PH)	Samples
2447231 proteinte 1,96 133,47 09 244701 AA4453310 proteinte 2,38 114,00 48 244703 AA4789879 proteinte Y 24,3 114,00 30 245768 AA5286 proteinte Y 24,6 423,6 17 245768 AA5080 Leali Y 24,6 423,6 17 139103 AA6723 proteinte Y 24,6 423,6 17 139104 AA46723 proteinte Y 24,6 423,6 17 13910 AA46723 proteinte Y 24,6 423,6 17 13910 AA67723 proteinte Y 124 44,3 17 11000665 AA427781 - Y 13,4 44,10 13 1100665 AA67781 - Y 13,4 44,10 13 1100665 AA67781 - Y 13,6 13,4					Bound		!	;	740 63	185 70	7.38	52.32	22.54	10.36	•
646077 AA186879 probable 7 245 714,00 48 284701 MA5440 probable Y 245 78.14 30 284703 MA5240 probable Y 246 78.16 30 285003 MA50820 Locall Y 246 44.38 17 285005 MS5261 Locall Y 246 42.36 17 133130 PA52897 probable Y 246 44.38 17 133130 PA52897 probable Y 124 45.10 13 1000865 AA21781 L Y 124 45.10 13 1000865 AA21781 L Y 124 45.10 13 1000865 AA21781 L Y 124 45.10 13 1000866 AA21781 L Y 12.44 46.10 13 1000866 AA21781 L Y 12.44		788180	AA453310	prostate		2 .	133.47	2	48.00		17.67	31.06	53.59	35.90	•
26A701 N64840 prosisis Y 241 73.10 30 245708 AA70336 prosisis NA 256 78.64 30 245708 AA50236 coal coal 22 24.05 78.64 30 281003 N65826 coal coal coal 22 22 22 130605 AA67235 prosisis Y 2.66 44.38 17 1100865 AA67725 prosisis Y 1.24 45.10 13 1100865 AA67726 prosisis Y 1.24 45.77 10 1100865 AA67727 prosisis Y 1.24 45.77 10 1100865 AA67727 prosisis Y 1.24 45.77 10 1100865 AA67727 prosisis Y 1.24 45.77 10 1100867 AA67724 prosisis Y 0.20 0.14 11.77 11.78 11.77 <		646037	AA196979	prostate		2.38	114.00	8	65.24	70.47		A.2.A.	41.24	19.50	•
450049 AAVTG1369 prosesse NIA 2.56 78.64 30 246768 NIGS26		284701	N64840	alatemo	>	177	73.10	8	31.27	50.07	0.40		45.42	20.45	•
246768 NESCRE CODI 0.21 22 281003 NEGERE - Coli 24.05 423.65 18 138606 FR3051 PR3387 Provide 7 2.66 17 138100 PR3287 Provide Y 2.64 17 13810 PR3287 Provide Y 1.24 46.10 13 170750 H20747 - Y 1.24 46.10 13 100066 AA67278 Done Y 1.24 46.10 13 100066 NB1003 - Y 1.24 46.10 13 100066 NB1003 - Y 1.24 46.10 13 202217 NB1003 - Y 0.50 6.84 9 202217 NB1003 - Y 0.50 6.84 9 202217 NB1003 - NA 1.81 12.19 1 20221 AA60478	, ,	450040	AA7m396	etatatva	¥	2.55	76.64	ន	38.52	11.57	\$4.04 1		3.44	63.61	•
24,050 Household 24,05 423.65 18 139130 R28397 Proteiale Y 2.66 44.38 17 139806 R28397 Proteiale Y 2.66 44.38 17 1000865 AA457235 Proteiale Y 1.48 25.46 17 1000865 AA457261 - - 0.12 12 12 1000865 AA42781 - - 2.07 19.56 9 1000865 AA677261 - - 0.01 0.12 10 1000866 Nig1003 - Y 0.20 0.64 9 200086 Nig1000 - - 0.50 0.84 9 21831 T72843 breast - 0.50 0.32 137 7 228359 AA00478 - - 0.00 0.44 4 4 4 4 4 4 4 4 4 4<		00000	***************************************		į	0.01	0.21	8	1.34	8.80	10.09	77.7		\$	•
13860G FASEST V 2.66 4.38 17 13860G FASEST Proteiale Y 2.64 44.10 13 13860G AAASTZSS proteiale Y 1.24 46.10 13 102066S AABTZTGS - 2.07 19.56 9 9 10506G Nalocas - 2.07 19.56 9 9 10506G Nalocas - 0.00 3.83 8 7 20527 Nalocas - 0.00 3.83 1 7 1 20527 AACOLAGSI - - 0.00 0.42 7 1 1 1 1 1 1 1	1 0	00/047	0070CN	• :		2	423.65	8	22.19	20.64	22.40	15.57	19.71	7 7	•
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153130 R28597 prosisis Y 1455 2240 153868 AA457255 prosisis Y 124 44:10 13 1600665 AA457255 prosisis 0.012 0.12 12 1600665 AA45725 prosisis 0.012 0.12 12 200066 Nations	~	136605	R35051		> -	8 :	00°#	: ţ	30.48	4:34	6.15	9.24	10.23	8	9
### 19868 AA45725 prosisis	•	£155	R26397	prostate	>	. \$	25.46	≥ 9		50.28	3.85	4.31	3.32	7.01	•
178753 H51649 bone 0.01 0.12 12 12 12 12 12 12 12	•	839668	AA457235	prostate		3	46.10	<u> </u>	3 8	3 3	200	10.01	17.09	10.66	•
51433 H20747 124 12.57 10 51433 H20747 12.4 12.57 10 306066 N91003 2.07 19.56 9 228215 R91386 0.06 0.42 7 228217 N78180 bone . 0.06 0.42 7 228218 AA004681 . 0.06 0.42 7 195034 R88764 . 0.10 0.60 0.42 243841 AA01178 . 0.10 0.60 0.14 243841 AA01178 .		179753	H51549	poue		<u>0</u>	0.12	5	9.28	8.5	***	6.14	4.27	15.03	•
61433 H20747 Y 0.80 6.84 9 300066 Ni91003 - 0.50 3.83 8 21631 T72543 breast 0.20 1.97 7 292217 N79180 bone - 0.06 0.42 7 428816 AA00481 - NIA 1.91 12.19 6 428816 AA00481 - NA 1.95 0.14 6 242841 AA01176 - AA01176 - 0.02 0.14 6 242841 AA01176 - AA01176 - 0.02 0.38 13 250374 AA01172 - AA01176 - 0.16 144.72 <td< th=""><th>: =</th><th>1030855</th><th>AA621761</th><th>•</th><td>></td><td>7</td><td>12,57</td><td>\$</td><td>7.99</td><td>19.6t</td><td>2 5</td><td>7 43</td><td>35.50</td><td>19.24</td><td>•</td></td<>	: =	1030855	AA621761	•	>	7	12,57	\$	7.99	19.6t	2 5	7 43	35.50	19.24	•
3000066 NINTOCK V 0.80 6.94 9 21831 77243 breast 0.50 3.83 8 228217 N79180 bone 0.42 7 225559 AA504784 - 0.06 0.42 7 422516 AA004681 - 0.00 0.42 7 422516 AA004681 - 0.00 0.00 0.14 6 422516 AA004681 - Maybe 1.26 0.14 6 422516 AA011176 - Maybe 1.55 0.14 6 245624 N40548 - Maybe 1.56 0.34 4 245621 N40548 - Maybe 1.56 0.34 4 245468 NA52103 prostate 1.56 0.34 4 4 256374 NY3522 prostate 1.59 13.52 10 1031539 AA000483 - 0.00 <	: 2	51433	H20747	•		2.07	19.56	a	6.78	6.32	0 1	1	3.65	6.31	•
195076 R91306 - 0.50 3.83 8 8 132517 N72543 breast 0.29 1.07 7 7 7 7 7 7 12543 breast 0.29 1.07 7 7 7 7 7 1 12553 breast 0.29 1.07 7 7 7 7 7 1 1255359 Ax504784 - 0.00 0.10 0.60 0.42 7 7 1.00 0.00 0.42 7 7 1.00 0.00 0.42 7 1.00 0.00 0.14 0.00 0.14 0.00 0.14 0.00 0.14 0.15 0.14 0.10 0.00 0.14 0.15 0.15 0.14 0.15 0.14 0.15 0.15 0.14 0.15 0.14 0.15 0.14 0.15 0.15 0.14 0.15 0.14 0.15 0.15 0.14 0.15 0.14 0.15 0.14	! !	2000	Notor		>	080	3 .9	o	13.54	16.10	S8: 7		1, 6,	6.41	•
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202217 N79180 bone i 0.06 0.42 7 225856 AA50484 - NAA 1.91 12.19 6 428816 AA004881 - 0.10 0.80 0.42 196024 R88764 - 0.10 0.10 0.80 0.14 196024 R88764 - 0.10 0.00 0.14 283701 AA011176 - 2.91 12.87 4 283203 H9888 - 2.91 12.87 4 283203 H9888 - 0.32 0.86 4 284582 NA22103 prositie 15.97 194.72 12 2263214 N31852 prositie 12.8 13.52 10 226321 N2986 prositie 12.8 13.52 10 226321 N2986 prositie 0.09 0.74 8 2642468 AA69689 - NAA 3.26 2.049 6 242458 AA69689 - NA 3.26 2.049 8611562 AA65487 bone Y 3.56 2.122 6 242436 AA004412 - 0.04 0.20 0.00 120139 R20913 - 0.04 0.20	•			,]		0.20	1.97	~	3.24	10.05	9 .80	7.18	9 1		• •
225277 N78180 Done 225859 AA50784 . NA 1.91 12.19 6 428816 AA00481 . 0.10 0.80 6 196034 R88764 . 0.10 0.00 0 242816 AA0041178 . 0.10 0.00 6 243641 N46548 . Maybe 1.55 6.34 4 243641 N46548 . 0.30 3.88 13 256374 N3162 prosisis 129 13.52 10 256374 N3162 prosisis 17.59 15.87 8 266275 AA60465 . NA 3.26 2.049 6 264276 AA64697 bone Y 3.59 21.22 6 256436 AA00412 . NA 3.26 2.049 6 256436 AA00412 . 0.04 0.20 6 256436 AA00412 . 0.04 0.20 6	L	2802	172343	Dieses	•,	3 5	0.42	7	3.84	10.11	7.14	10.04	4.07	: t	•
### ### ### ### ### ### ### ### ### ##	•	282217	N79180	Poop	•	3 6	4 49	· •c	6.28	4.64	5.38	5.13	5.23) e :	•
42816 AAO04681	_	825859	AA504784	•	₹			· «	45.8	8.44	8.60	4.79	4.07	3.73	۰ و
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359701 AA011176 2.91 12.57 4 243641 N46548 - Asybe 1.55 6.34 4 243641 N46548 - 0.22 0.86 4 287035 H68688 - 0.22 0.86 4 256374 N31652 prostate 1.59 19.72 12 256374 N31652 prostate 1.29 13.52 10 256374 N31652 prostate 1.29 13.52 10 256374 N31652 prostate 1.29 13.52 10 256374 N31652 prostate 1.29 15.87 10 163620 H67712 - 0.19 1.82 10 163620 H6177 - 1.75 15.29 10 163620 H6177 - 1.75 11.25 10 163620 H6177 - 1.75 11.25 10 163621 H72336		195034	R86764			70.0	<u>•</u>	•	3.21	3.87	3.89	8.96	3.63	3.06	Đ
243641 N46546 - Maybe 1.55 0.34 287035 H98689 - 0.22 0.86 4 287035 H98689 - 0.30 3.88 13 287036 NA3103 prostate 1.29 19.72 12 259374 N31652 prostate 1.29 13.52 10 193923 R83407 brain Y 17.59 158.73 9 1031590 AA60443 bone 0.09 0.74 8 403463 AA6986 prostate N/A 0.37 3.71 6 422468 AA64597 bone Y 3.59 6 428436 AA644597 bone Y 3.59 6 428436 A	_	359701	AA011176	•		2.91	12.67	• •	3.47	88.5	4.16	3.84	3.52	3.85	•
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37990 R61372 lung 0.06 0.74 8 1031599 AA600483 - 0.27 2.07 8 86160 T72336 bone 1,71 11.25 7 208211 NZ9966 prostate 0.57 3.71 6 442468 AA699895 - N/A 0.38 2.39 6 866276 AAA64597 bone Y 3.56 21.22 6 428436 AAA64597 bone Y 3.59 21.22 6 428436 AAA64437 bone Y 3.59 21.22 6 130136 R20613 - 0.04 0.20 6	_	193923	R83407	brain	> -	17,59	158.73	œ	10.87	9.6	5 6	60.	4.81	35.0	6
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86160 T72336 bone 1,71 11.25 7 200211 N29666 prosisio 0.67 3,71 6 462468 AA690895 NIA 0.38 2.39 6 806276 AA653160 NIA 3,26 20.49 6 811562 AAA54597 bone Y 3,59 21.22 6 428436 AA00412 0.65 3,12 6 130136 R20813 0.04 0.20 6	_	1031599	AA609485	•		0.27	2.07	ω.	5.69	10.96	9.75	<u> </u>	7.88	4,49	10
208211 N29986 prosisio 0.67 3.71 6 422468 AA690895 . N/A 0.38 2.39 6 806276 AA653160 . N/A 3.26 20.49 6 811582 AAA54597 bone Y 3.59 21.22 6 428436 AA00412 . 0.53 3.12 6 130136 R20813 . 0.04 0.20 6		86160	172336	e coq		1.7	11.25	۲	7.39	5.16	3 1	5.5	55.4	6.18	40
482468 AA690895 . NVA 0.38 2.39 6 866276 AA653160 . NVA 3.26 20.49 6 811562 AAA54597 bone Y 3.59 21.22 6 428436 AA00412 . 0.53 3.12 6 130136 R20813 . 0.04 0.20 6		268211	N29986	prostate		0.57	3.71	89	8.59	3.86	13.77	21.2	3 7	3.81	10
866276 AA663160 - NA 3.26 20.49 6 811582 AAA64597 bone Y 3.59 21.22 6 428436 AA00412 - 0.53 3.12 6 130136 R20813 - 0.04 0.20 6		442448	AABOOBOR		N/A	920	2.39	90	5.95	10.82	5.45 8	10.02	<u> </u>		· •
811582 AAA54597 bone Y 3.59 21.22 6 428436 AA004412 . 0.53 3.12 6 130136 R20813 . 0.04 0.20 6	٠.	******	A A B 621 4 B	• 1	A	3.26	20.49	9	1.91	3.57	4.27	14.38	3 (3	,	· •
428436 AA004412 . 0.65 3.12 6 130136 R20613 . 0.04 0.20 6			2012000	. ;	>	9,6	21.22	ထ	9.12	4.48	4.7	9	2.69	6 1	, u
428436 AA004412	•	700110	/AC+C+47		-			• •	1.13	4.65	£.73	5.7.	11.11	8.	D 1
130136 R20613 . U.D. U.A. C. C.	10	428436	AA004412	•		3 3	. c	• ¢	7.23	6.80	5.30	5.38	9 .89	2.08 80	.
	•	130136	R20613			\$ 3	9 6) (2 80	4.97	4.10	7.9	3.72	11.48	ю .
W81432 heart 0.04 0.24 0	~	347586	W81432	hear		3	770	o (8	0.83	11.93	6.34	3.73	¥Ď
38 795729 AA480291 unspecified 0.02 0.11 6 5.09	•	795729	AA460291	petipedeun		0.02	0.11	w	AO.C	2	}				
									•						

Table 2-2: T3NO vs. BPH: Gene uplegulated 5 fold of higher in at least 2 tumor samples or 3 fold or higher in at least 3 tumor Samples.

	!	Gen Bank		Secreted/Me			Fold(AveTumor/A Fold(MPM120/A Fold(MPM121/A Fold(MPM124/A Fold(MPM122/A Fold(MPM125/A #ofTumors Fold(MPM125/A #ofTumors Pold(MPM125/A #ofTumors Fold(MPM125/A #ofTumors Pold(MPM125/A #ofTumors Fold(MPM125/A #ofTumors Fold(M	Fold(MPM120/A	Fold(MPM121/A	Fold(MPM124/A	Fold(MPM122/A	Fold(MPM123/A	Fold(MPM125/A i	# of Tumora Samples
ode	Clone 1D	Accession	Distribution	mbrane	AveBPH	AveTumor	veBPH)	veBPH)	√•8 ₽₩)	(LL)			4.56	10
		Number		Bonua	į		*	3.24	6.98	2.85	6.31	5		
8	825785	AA505117	Ovary		2.	i i	, u	2.31	3.63	6.60	3.61	13.00	0.50	
\$	49858	H29290	brain		S	3	· •	2.79	3.52	3.68	4.21	8 00:	Ż.	
=	417075	W87823	•		0.83	4.38	6 (75.6	10.29	3.87	2.38	7,56	4.13	Ω 1
4	840751	AA486071	•	≻	ŗ.	00 00	o 1		6.61	5.92	8.08	2.79	33	n 1
1	241821	H93381	•	¥	8.	5.43	.	2 2	8	4.57	7.80	4.62	2.43	o 1
: 3	29063	R40970	prostate		3 .	8.52	NO.	07.4	3 2	68	4.48	8.48	5.11	1 0
; ;	124128	801348		>	1.25	6.16	80	2.53	5 6	8	11.43	4.02	4.20	1 0
2 :	20000	02030	Anmy		4.70	23.13	NO.	2.33		3 5	ă	3.53	2.83	1 0
\$	70007	2000	i de la constante de la consta		0.29	54.	ĸ	5.02	7.58	9 1		3.82	5.12	١n
Ş	201	K2828/	5		3.81	18.41	ĸ	2.38	4.28	3.75	, F	6.37	4.56	w
\$	28//82	W62244			0	90:0	vo	0.79	8.80	9 .4		7,	4.47	10
4	271141	X34482	•		, c	9.60	vo	1.68	4.13	3.48	9 C.	200	3.63	10
8	1240414	AA788780	• ;		5	15.28	10	9.25	5.31	0.40	80°C	7	10.4	w
2	307249	N83438	testis		2 .	9 0	•	4.82	5.19	6.45	4.67	<u> </u>	2	10
8	763849	AA443688	OVBIY		2 6	9 6	. •	3.68	3.50	2.84	3.83	9 ;	200	, v a
ន	897720	AA598982	brain		0.32	8 9	• •	3.87	98.9	3.48	5.50	ă,	2	, w
3	283070	N51297	testis		8	21.0	• •	8.18	3.28	3.22	9	0.56	9.0) W
8	265005	N21081	akļu		90.0	0.20	•	. e	7.98	3.20	3.85	0.24	06.5	
8	343987	W70234	poue		8 .57	35.37	• •	 	7.08	3.45	3.00	5.14	3.31	.
150	416655	W87281	•	>	0. 84.	4.88	•	2 0	463	2.64	4.	5.85	3.05	6 1
3	842848	AA486281	Brug	> -	1.15	4.62	•		183	4.50	3.68	4.50	2.31	o 1
2	40690	R55882	brain		90.0	0.18	•	3.23		3.16	3.29	3.49	3 .0	ю .
8	359661	AA011096	prostate		23.32	84.82	•	0.4) 	3.88	4.66	4.80	0.80	n '
	126230	R06307	prairy		0.03	0.10	e -	3.18	326	3.51	3.03	3.10	1.45	.
2	375682	AA032221	prostate	>	15.66	45.43	eo '	3.03	9	50.34	105.24	0.80	19.00	•
2	1468260	-	•	ΥN	0.65	22.38	ಕ್ಷ :	0.00	78 11	29.56	0.86	23.62	1.26	→ ·
3	743828		poue	₹	0.58	12.90	23	29.78	1873	3.22	1.15	2.77	3.51	₩ .
. 2	1390584	AA843592	prostate		<u>7</u> .	33.09	8	48.03	5 to 0	4	57.03	15.40	1.72	•
: 2	41647	R52794	Drei		0,40	6.34	5	6.8	5 5	98	8.52	48.21	5.35	+
2	744944	AA625890	prostate		5. 8.	22.42	12	2.70	2 7	3.20	9.26	2.32	1.91	→
: \$	1034473	AA770728	prostate	\	0.13	1.36	=	22.48	£3:74 10.45	13.83	13.97	6.5	6 .	•
2	487165	AA045074	rojos		0.0	0.0	æ	13.49	54 to	88	8.31	4.84	2.76	→
2	1456424	AA862992	breast		90.0	0.69	ω	50.01	3 5	230	24.39	4.57	6.15	•
: =	360155	AA013260			0.01	0.11	ထ	10.07	5.5	2 6	10.46	<u>7</u> .	2.63	~
3	144740	R76229	prostate		0.19	1.27	_	7.58	DC:01	7 3 4	2.85	10.25	15.67	₹
: #	24915	R39066			0.83	5.97	9	3.07	2.31) Y	9.80	2.35	3.29	→
? ;	ROTTER	4446678		>	9.70	60.24	\$	9.37	12.88	?		1.74	10,42	•
٤;	200	-	• Deline	•	11.61	70.00	စ	6.66	5.08 8.08	rr.01	• E		2.67	•
2	7/1001	•			800	80 80	9	8.95	11.77	5.12	20.0	90.	3.10	4
2	782383	-	•		2	70.0	\$	16.58	0.83	4,9	4.81	8 8	74.6	•
=	1031113	•	•	;			¥C	2.60	6.47	9.51	6.43 5.43	20.0		•
28	461541	AA705258		Š	5 6	3 5	u u	3.93	6.17	7.31	9.03	2.28	107	• •
2	491706	A115259	breast		6.79		o ur	6.20	9.84	6.65	1.30	0.26	9	•
2	787938	AA452278	prostate	>	1.76	0.0	,							
								7						

Table 2-2: T3NO vs. BPH: Gene upregulated 5 fold of higher in at least 2 fumor samples or 3 fold or higher in at least 3 fumor Samples.

5 5.74 5 7.74 6 7.74 7 7.74 7 7.74 7 7.74 7 7.74 7 7.74 7 7.75 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7	Gen Bank Accession		₹ :		Secreted/Me mbrane	AveBPH	AveTumor	Fold(AveTumor/A Fold(MPM120/A Fold(MPM121/A Fold(MPM122/A Fold(MPM123/A Fold(MPM125/A # ofTumors	Fold(MPM120/A	Fold(MPM121/A	Fold(MPM124/A	Fold(MPM122/A veBPh)	Fold(MPM123/A veBPH)	Fold(MPM125/A veBPH)	# of Tumors Samples
5 574 889 492 206 177 6 108 359 1096 825 064 177 6 310 354 625 246 277 143 625 5 324 626 274 828 276 177 6 139 430 463 274 328 107 6 623 1070 441 526 270 117 6 623 1070 443 828 270 117 6 623 1070 441 526 117 6 625 650 839 328 338 146 6 626 630 177 142 110 142 6 627 626 839 339 148 147 6 627 626 833 101 150 238 6 627 626 839	Distribution Bound	Distribution Bound	Bound									5	46.8	9.17	· 🕶
5 302 3138 482 220 0.14 6 319 354 626 226 0.04 6 319 334 626 226 0.04 6 319 430 274 928 0.04 6 189 473 743 326 3.06 6 189 473 743 326 3.06 6 48 483 274 928 442 6 48 483 274 928 119 6 48 483 328 751 119 6 48 483 328 751 119 7 48 483 373 441 119 8 560 344 528 173 118 8 560 347 373 441 141 8 16 242 171 148 141 8 1	754034 AA479058 . 0.02	•	0.02	0.02	0.02		0.10	NO.	5.74	6.97	3	70'1	7.	328	•
6 1186 3.59 10.39 85.22 0.024 6 3.19 3.44 0.63 4.65 2.50 11.89 0.024 6 3.19 4.80 2.74 9.28 2.70 157 6 1.92 9.73 7.43 9.28 2.70 157 6 5.23 10.10 4.41 9.28 2.70 150 6 6.25 6.50 9.73 7.51 1.16 1.16 7 6.26 5.50 9.33 1.01 1.85 2.70 1.80 8 6.27 10.70 4.41 9.28 1.16	AARBOT77 bons	*		900	\$0.0		91.0	ĸ	3.02	13.78	4,92	2.90		3 5	-
5 3.22 3.11 2.03 1139 5.25 6 3.19 3.24 6.86 2.56 2.70 6 1.92 4.80 2.74 9.28 4.42 6 5.23 10.10 4.41 6.28 1.37 6 5.22 10.10 4.41 6.28 1.39 6 5.22 10.10 4.41 6.28 1.39 6 5.22 10.10 4.41 6.28 1.39 7 4.85 3.86 3.87 9.59 1.39 6 4.85 3.87 9.59 1.39 1.35 6 4.06 2.42 10.28 1.41 1.41 1.43 1.44 7 4.06 2.47 3.73 3.49 4.41 1.41 1.44 1.44 1.44 1.44 1.44 1.44 1.44 1.44 1.44 1.44 1.44 1.44 1.44 1.44 1.44 1.44	T63389 hoos	2 60		0.02	0.02		90.0	ĸo	1.68	3.59	10.99	6.52	* 000		•
5 319 394 6.86 2.76 1.71 1.17 6 1.92 9.73 7.83 3.66 3.71 1.17 1.17 6 1.92 9.73 7.83 3.65 3.88 1.16 4.42 5.89 1.16 4.42 5.89 1.16 4.42 5.89 1.16 4.42 5.89 1.16 4.42 5.89 1.16 4.42 5.89 1.16 4.42 5.89 1.16 4.42 5.89 1.16 4.42 5.89 1.16 1.18 4.42 1.10 1.18 4.42 1.10 1.18 4.42 1.10 1.18 4.42 1.10 1.18 4.42 1.10 1.18 4.42 1.10 1.10 4.42 4.42 1.10 1.18 4.42 4.42 1.10 1.13 4.41 4.41 4.41 4.41 4.42 4.42 4.41 4.41 4.41 4.41 4.41 4.42 4.42 4.42 4.42	STORY CONTRACTOR			3.5	3.53		17.34	wo.	3.22	3.11	2.03	11.89	6.25	/A.2	• •
5 3.44 0.53 4.85 7.12 1.37 6 1.199 9.73 7.83 7.12 4.22 4.22 6 1.29 9.73 7.83 3.56 1.16 5.28 3.46 6 5.20 10.10 4.41 5.26 3.39 1.16 1.30 3.66 3.66 3.06 1.16 1.29 4.81 1.20 3.28 3.28 3.39 1.16 1.20 1.20 1.20 3.39 1.16 2.28 3.69 2.28 3.39 1.18 2.28 3.28 1.20 1.63 2.28 3.60 1.23 1.43	174302 (1944)	170	100				2.17	ĸ	3.19	3.94	6 .56	2.56	2.5	2 1	• •
5 1.59 4.80 274 9.29 4.42 5 5.25 1.010 4.41 6.26 1.16 5 5.25 10.10 4.41 6.26 1.16 5 5.60 10.81 5.89 3.36 1.16 6 4.62 6.50 0.33 1.01 1.85 6 4.64 2.42 10.26 1.76 4.81 6 4.64 2.47 3.73 6.40 6.89 7 1.85 2.47 3.73 6.40 6.89 8 2.47 3.73 6.40 6.89 2.89 1.36 2.89 1 1.85 2.47 3.73 1.00 6.89 2.89 6.89 3.89 6.70 2.89 6.70 2.89 6.70 2.89 6.70 2.89 6.70 2.89 6.70 2.89 6.71 6.70 6.70 2.89 6.71 6.70 6.70 6.70 6.70	Office of the second control of the second c	0.02	0.02			o	8	vo	3.4	0.53	3 .	7.12	/c.T	3 ;	• •
5 1.92 97.3 7.83 3.66 3.86 6 5.23 10.10 4.41 5.29 1.19 6 5.23 10.10 6.44 5.29 1.19 6 5.52 6.50 6.33 7.51 1.30 6 4.85 3.66 3.87 9.59 2.38 7 4.85 3.66 3.87 9.59 2.38 6 4.85 3.66 2.42 1.02 1.85 3.38 7 1.87 3.44 4.51 1.56 3.38 1.38 3.38 4 1.87 3.44 7.71 1.36 3.38 1.38 3.38 4 5.48 6.46 6.39 7.71 1.36 3.89 5.48 5.99 2.50 7.73 6.18 3.60 4.67 6.87 3.20 3.53 4.59 7.74 6.18 6.87 2.48 4.09 2.74 <		500	900				24	ĸ	1.89	4.80	2.74	9.28	1.42	C (• •
5 10,10 441 5,28 1,16 6 0,48 483 6,38 7,51 1,30 6 5,52 5,50 10,31 1,53 1,30 6 5,52 5,50 10,31 1,53 1,39 7 1,68 3,68 3,87 9,69 2,38 4 1,66 2,47 1,026 1,78 4,81 5 1,69 2,47 1,026 1,176 4,81 4 5,34 6,46 6,39 1,35 3,86 1,36 3,86 4 5,16 4,37 3,15 6,13 1,37 3,86 1,17 4,81 1,18 3,86 1,18 3,86 1,18 4,81 4,	AACOO 1 1000 1000 1000 1000 1000 1000 1000	0.27	0.27				1.28	w	1.92	9.73	7.83	3.56	3.08	52 1	• •
6 0.48 4.83 7.51 1.30 6 5.56 6.39 7.51 1.30 6 4.85 5.86 3.33 1.01 1.85 6 4.85 3.66 3.87 9.59 2.38 1.38 6 4.85 3.66 3.87 9.59 2.38 1.81 7 1.85 2.47 1.02 1.76 4.81 1.65 6 2.47 5.43 7.71 1.35 2.38 1.86 2.86 2.89 2.89 2.89 2.89 2.89 2.89 2.89 2.89 2.89 2.89 2.89 2.89 2.89 2.89 2.89 2.89 2.89 2.89 4.87 4.87 4.87 4.87 4.87 4.87 4.89 4.89 4.89 4.89 4.89 4.89 4.89 4.89 4.89 4.89 4.89 4.89 4.89 4.89 4.89 4.89 4.89 4.89 4.89	580	0.00	590			•	5.42	ĸ	5.23	10.10	4.41	5.26	1.16	2.07	•
5 3.60 10.81 5.89 3.38 1.85 5 5.52 6.50 9.33 1.01 1.63 6 4.86 3.66 9.33 1.01 1.63 6 4.04 2.47 10.26 1.76 4.81 6 1.86 2.47 3.73 5.40 9.80 7 1.86 2.47 3.73 5.40 9.80 6 1.86 2.47 3.73 5.40 9.80 7 2.17 3.26 4.81 3.80 2.38 8 5.34 4.77 3.18 3.74 4.87 8 5.48 5.80 2.80 2.80 2.80 8 5.34 4.59 7.74 4.87 8 5.79 4.89 3.74 4.87 8 6.87 3.20 2.70 4.77 3.89 8 6.10 3.24 4.77 3.89 4.72			200				10	NO.	0.48	4.83	6.38	7.51	8.	7.00	• •
5 5.52 6.50 9.33 1,01 1,63 6 4,88 3.66 3.87 9.59 2.38 6 4,88 3.66 3.87 9.59 2.38 7 1,18 2.47 1,35 5.49 4.81 4 2,17 6.46 6.49 1,35 5.49 6.80 4 2,17 5.43 4.73 1,005 2.98 6.80 4 5,19 6.49 2.50 2.50 9.71 1,81 5 6.19 4,37 3.26 4.59 7.43 6.18 5.28 4 5,48 5,59 2.50 3.29 4.69 4.67 4 1,27 3.20 5.81 2.50 1,75 6.18 4 2,18 5.22 3.69 5.17 8.99 4.72 4.23 4 1,27 3.01 3.89 4.72 4.23 4.73 4.43 4 <th>170</th> <th>170</th> <th>170</th> <th></th> <th></th> <th></th> <td>ā</td> <td>ĸ</td> <td>3.60</td> <td>10.81</td> <td>5.89</td> <td>3.38</td> <td>1.85</td> <td>2.40</td> <td>• •</td>	170	170	170				ā	ĸ	3.60	10.81	5.89	3.38	1.85	2.40	• •
5 486 368 387 9.99 2.38 6 4,04 2.47 3.73 6.40 6.89 2.88 6 6.186 2.47 3.73 6.40 6.89 1.76 4.81 6 6.18 6.46 6.39 1.35 6.02 6.02 6 6.19 6.40 6.39 1.35 6.02 6.02 6 6.19 6.43 7.71 1.35 3.89 6.29 6.02 6 6.48 6.59 2.50 3.20 3.20 3.20 3.20 4.71 1.81 6 6.48 3.20 5.81 2.60 1.75 4.23 4.72 4.87 6 6.89 3.21 2.74 3.89 4.89 4.87 4.87 4.87 4.87 4.89 4.89 4.89 4.89 4.89 4.89 4.89 4.89 4.89 4.89 4.41 4.41 4.41 4.41 4.41 <th>2000 COOPER</th> <th>100 CT</th> <th>4 C C C C C C C C C C C C C C C C C C C</th> <th></th> <th></th> <th>. K</th> <td>8</td> <td>vo</td> <td>5.52</td> <td>5.50</td> <td>9.33</td> <td>1.01</td> <td>3.</td> <td>\$.</td> <td>• •</td>	2000 COOPER	100 CT	4 C C C C C C C C C C C C C C C C C C C			. K	8	vo	5.52	5.50	9.33	1.01	3.	\$.	• •
5 4,04 2,42 10,26 1,78 4,81 6 1,86 2,47 3,73 5,40 6,60 6 6,46 6,39 1,35 5,02 6 1,17 5,43 7,71 1,35 3,96 6 5,19 4,37 3,15 9,71 1,81 3,96 6 5,19 4,37 3,15 9,71 1,81 3,96 4,97 3,96 2,99 4,97 3,96 4,97 3,96 4,97 4,97 4,97 3,96 2,74 4,97 4,97 4,97 4,97 4,97 4,97 4,97 4,97 4,97 4,97 4,97 4,97 4,97 4,97 4,97 4,97 4,97 4,98 4,97 4,97 4,98 4,97 4,47 5,99 5,17 8,99 4,47 5,19 6,19 6,17 4,99 4,47 5,19 6,17 4,41 4,47 5,19 4,47 5,19 4,47	7.4.7	prosterio	247	7.4.7		÷	22	wo	4.85	3.66	3.87	9.50	2.38	2.8/	• •
4 1.85 2.47 3,73 5.40 9,80 4 5.34 6.46 6.39 1.35 5,02 4 5.17 5.43 7.71 1.35 3.99 4 5.19 4.37 3.15 9.71 1.81 3.99 4 5.19 4.37 3.15 9.71 1.81 3.99 4 5.48 5.99 2.50 3.28 4.51 1.74 1.81 1.75 4 5.48 5.99 2.50 3.29 4.57 4.51 1.75 4.67 4.67 4.67 4.67 4.67 4.67 4.67 4.69 4.77 4.69 6.19 6.17 6.19 6.19 6.17 6.19 6.17 6.19 6.17 6.18 6.17 4.71 4.71 4.71 4.71 4.71 4.71 4.71 4.71 4.71 4.71 4.71 4.71 4.71 4.71 4.71 4.71 4.71 4.71	2021/VV	980 	200 of 100 of 10	96		¥	22	NO.	4.04	2.42	10.26	1.76	4.81	3.75	• •
4 5.34 6.46 6.30 1.35 5.02 4 2.17 5.43 7.71 1.35 3.96 4 5.19 4.37 3.15 9.71 1.81 5 1.97 3.35 4.29 10.05 2.98 6 1.97 3.35 4.59 7.43 6.18 6 6.87 3.20 5.81 2.50 1.76 6 8.77 3.29 7.43 6.18 6 8.87 3.20 5.81 2.50 1.75 6 8.87 3.20 5.81 2.50 1.75 6 8.88 2.46 4.06 2.17 8.08 6 8.99 3.21 2.71 1.052 0.89 6 8.99 3.24 4.06 2.17 8.08 6 8.11 2.26 3.24 4.41 1.37 4.41 6 8.12 8.24 4.47 6.19	ACC.	700	760	đ		7	- 10	*	1.85	2.47	3.73	5.40	9.60	* 1	• •
4 2.17 5.43 7.71 1.35 3.86 4 5.19 4.37 3.15 9.71 1.81 4 5.19 4.37 3.15 9.71 1.81 4 5.19 4.37 3.15 9.71 1.81 4 5.86 6.89 2.50 3.26 4.73 6.18 4 2.86 6.89 3.63 4.72 4.28 4 2.89 3.63 4.72 4.29 1.75 4 2.89 3.21 2.71 1.052 0.69 4 6.17 5.02 3.89 2.17 8.08 4 6.17 5.02 3.89 2.74 2.97 4 6.17 5.02 3.89 2.74 4.41 5 5.03 4.77 3.89 6.17 4 5.19 5.74 4.41 3.29 4 5.19 5.74 4.41 3.29 4 5.19 5.74 4.41 3.29 4 5.19 5.29 3.74 4.30 4 4.17 5.19 3.29 5.10 4 4.17 5.19 3.29 4.41			240			20.		*	5.34	6.46	6.30	1.35	5.02	r.,	• •
4 3.25 3.54 4.23 10.05 2.88 5.19 4.37 3.15 8.71 1.81 6.19 4.37 3.15 8.71 1.81 6.19 5.89 2.50 3.28 4.67 7.43 6.18 6.87 3.20 4.59 7.43 6.18 6.18 6.19 6.17 6.18 6.87 3.20 5.81 2.50 1.75 6.18 6.17 6.18 6.19 6.17 6.18 6.19 6.17 6.18 6.19 6.17 6.28 6.09 2.17 6.18 6.18 6.19 6.17 6.18 6.19 6.17 6.19 6.17 6.19 6.17 6.19 6.17 6.19 6.17 6.19 6.17 6.19 6.17 6.19 6.17 6.19 6.17 6.19 6.17 6.11 6.11 6.11 6.11 6.11 6.11 6.11 6.11 6.11 6.11 6.11 6.11	- Coord	***				Č		- 4	2.17	5.43	1.71	1.35	3.88 3.88	5.83 62	•
4 5.19 4.37 3.15 9.71 1.81 5.48 5.99 2.50 3.26 4.67 6.87 3.20 5.81 2.50 1.75 4 2.86 6.89 3.63 4.72 4.28 4 2.86 6.89 3.63 4.72 4.28 4 3.89 3.21 2.71 10.62 0.69 4 1.27 3.01 3.96 5.13 8.53 4 6.11 5.02 3.77 10.62 0.69 4 6.13 3.29 4.77 8.69 6.17 2.97 4 6.13 3.69 5.13 8.53 4.41 2.97 4 6.14 5.02 4.77 3.69 6.17 2.97 4 6.19 5.74 4.41 1.37 4.41 2.97 4 6.10 6.74 4.47 5.19 5.29 4.40 4 4.67 </td <th>AA004828 unspecified</th> <th></th> <th></th> <th>•</th> <th></th> <th>9</th> <td>. *</td> <td>• •</td> <td>3.25</td> <td>3.54</td> <td>4.23</td> <td>10.05</td> <td>2.98</td> <td>1.97</td> <td>•</td>	AA004828 unspecified			•		9	. *	• •	3.25	3.54	4.23	10.05	2.98	1.97	•
4 548 5.99 2.50 3.26 467 4 1.97 3.35 4.59 7.43 6.18 4 6.87 3.20 5.81 2.50 1.75 4 2.86 6.89 3.63 4.72 4.28 4 3.89 2.48 4.06 2.17 8.08 4 3.89 3.21 2.71 10.52 0.69 4 1.27 3.01 3.86 5.13 8.53 4 6.11 5.02 3.69 2.74 2.97 4 6.13 3.89 2.74 4.41 5.19 5.19 3.29 7.01 4 5.19 5.19 3.29 4.60 4 5.19 5.19 3.29 4.60 4 5.19 5.19 3.29 4.60 4 5.19 5.19 3.29 4.60 4 4.67 5.61 2.56 4.74	A1003625 . N.A 4:42	NA A.42	77.4 VN	7 66		2	. כ	• •	5.18	4.37	3.15	9.71	1.81	3.5	•
4 1.97 3.35 4.59 7.43 6.18 4 6.87 3.20 5.81 2.50 1.75 4 2.86 6.88 3.63 4.72 4.28 4 3.89 2.47 4.28 6.89 4 3.89 2.17 8.08 4 6.17 3.01 3.89 2.74 2.87 4 6.17 3.01 3.89 2.74 2.87 4 6.17 3.01 3.89 2.74 2.87 4 6.17 3.01 3.89 6.13 8.53 4 6.17 3.01 3.89 6.17 2.87 4 5.06 5.74 3.81 2.10 3.39 4.41 4 6.40 3.53 3.79 6.85 1.28 4.60 4 4.67 5.61 2.59 3.74 4.10 4.10 4 4.67 5.61 2.65 4.74	H22854 brain 1.94	Drain 1.96	AU	7 6		2 6		•	84.8	5.99	2.50	3.26	4.67	2.80	.
6.87 3.20 5.81 2.50 1.75 4 2.86 6.88 3.63 4.72 4.28 4 2.86 6.88 3.63 4.72 4.28 4 3.89 2.48 4.06 2.17 8.08 4 1.27 3.01 3.86 5.13 8.53 4 6.11 5.02 3.89 2.74 2.87 4 5.08 5.00 4.77 3.89 6.17 5.19 5.00 4.77 3.89 6.17 4 5.19 3.74 4.41 4.41 5.19 5.74 4.77 3.89 4.60 4 2.57 4.47 5.19 3.29 4.60 4 2.03 4.18 2.88 3.29 4.60 4 4.67 3.33 3.82 2.44 4.10 4 4.67 3.53 3.82 2.44 4.10 4 4.16	AAA79484 breast Y 19.01	Dreast Y 10.00	10.01	15:01		3 6	۰.	•	1.97	3.35	4.59	7.43	6.18	1.26	→ ·
4 2.86 6.88 3.63 4,72 4.28 4 3.68 2.48 4.06 2.17 8.08 4 3.89 3.21 2.17 10.52 0.69 4 1.27 3.01 3.86 5.13 8.53 4 6.11 5.02 3.89 2.74 2.87 5 5.06 5.00 4.77 3.69 6.17 6 5.03 3.34 1.44 1.37 4.41 6.19 5.00 4.77 3.69 6.17 4 5.19 2.74 2.97 6.17 4 2.57 4.47 5.19 3.29 4.60 4 2.03 4.18 2.58 3.29 4.60 4 2.03 4.18 2.58 3.24 4.10 4 4.67 5.61 2.58 2.44 4.10 4 4.67 5.61 2.58 2.44 4.10	H66840 colon Maybe 0.23	colon Maybe U.13	Maybe	200) F		•	6.87	3.20	5.81	2.50	1.75	4.46	•
4 3.68 2.46 4.06 2.17 8.08 4 3.89 3.21 2.71 10.52 0.69 4 1.27 3.01 3.96 5.13 8.53 4 6.11 5.02 3.69 2.74 2.97 4 2.08 5.00 4.77 3.69 6.17 4 2.08 5.00 4.77 3.69 6.17 4 2.08 5.00 4.77 3.69 6.17 4 2.09 5.74 3.81 2.10 5.38 4 2.57 4.47 5.19 5.20 7.01 4 2.57 4.47 5.19 5.26 7.01 4 2.53 4.18 2.58 3.38 4.60 4 2.34 3.33 3.82 2.44 4.10 4 4.67 5.69 3.38 4.60 4.10 4 4.67 5.69 3.36 5.17 4.30 4 4.17 5.69 5.17 4.80 4.57	AA443080 bone bone 0.04	Pool Transfer Transfe	20:0	700	•	. 6	٠ «		2.86	6.98	3.63	4.72	4.28	2.07	•
4 3.89 3.21 2.71 10.52 0.69 4 1.27 3.01 3.89 5.13 8.53 4 6.17 5.02 3.69 2.74 2.97 4 2.08 5.00 4.77 3.69 6.17 4 2.08 5.00 4.77 3.69 6.17 4 2.57 4.47 5.19 2.10 3.38 4 2.57 4.47 5.19 3.20 7.01 4 2.57 4.47 5.19 3.20 7.01 4 2.93 4.18 2.58 3.38 4.60 4 2.94 3.35 3.52 2.44 4.10 4 2.94 3.33 3.82 2.44 4.10 4 4.67 5.61 2.56 4.74 3.85 4 4.67 5.61 2.55 4.74 3.85 4 4.16 2.79 3.26 4.74 4.30 4 4.16 2.79 3.26 5.74 4.30 4 4.16 2.79 3.26 5.74 4.30 4 4.16 2.79 3.26 5.74 4.30 <		akin maybe	maybe - 172	1.72		0.7	,	*	3.68	2.48	4.06	2.17	80.8	1 (1)	• •
4 1.27 3.04 3.89 5.13 8.55 4 6.11 5.02 3.69 2.74 2.97 4 2.08 5.00 4.77 3.69 6.17 4 5.19 5.74 3.81 2.10 3.38 4 2.57 4.47 5.19 3.20 7.01 4 2.57 4.47 5.19 3.29 7.01 4 2.93 4.18 2.88 3.38 4.60 4 2.94 3.35 3.82 2.44 4.10 4 4.67 5.61 2.56 4.74 3.85 4 4.67 5.61 2.56 4.74 3.85 4 4.67 5.61 2.55 4.74 3.85 4 4.67 5.61 2.55 4.74 3.85 4 4.16 2.79 3.26 5.17 4.30 4 4.16 2.79 3.62 5.06 3.32 4 4.16 2.79 3.81 6.06 3.32 4 4.17 4.61 5.01 4.80 4.57 4 4.61 5.01 4.80 4.57 4	11,5000 DOING 13,58	13.58	13.56	13.56		54.73	•	4	3.89	3.21	2.71	10.52	0.69	3.20	• •
4 6.11 5.02 3.69 2.74 2.89 5.17 2.81 2.74 2.81 2.74 4.41 4.42 4.42 4.43 4.46 4.40 4.	44702048 . 1.44	1,44	4).	<u>‡.</u>		#C	, _	•	1.27	3.01	3.96	9.13	3 6	<u> </u>	- ◀
4 2.08 5.00 4.77 3.34 1.44 1.37 4.41 4 5.19 5.74 3.81 2.10 3.38 4 2.57 4.47 5.19 3.29 7.01 4 2.83 4.18 2.58 3.38 4.80 4 2.83 4.18 2.58 3.38 4.80 4 2.84 3.33 3.82 2.44 4.10 4 4.67 5.81 2.55 4.74 3.85 4 4.67 5.81 2.55 4.74 3.85 4 4.67 5.15 3.36 5.17 4.30 4 4.67 5.16 3.36 5.17 4.30 4 4.16 2.79 3.62 5.06 3.32 4 4.17 6.29 3.11 1.20 3.29 4 1.21 4.61 5.01 4.80 4.57 4 5.81 3.02 3.16 2.19 2.78 5 5 5 3.77 3.86 1.82 5.82 3 3 3 3 3 3 3 4 5 4 4 4	AAA09106 hypothalamus 1.74	hypothalamus 1.74	1.74	1.74		9.0	¥	4	6.11	5.02	3.60	2.7	, t e	1.75	•
4 9.05 3.34 1.44 3.81 2.10 3.38 4 2.57 4.47 5.19 3.29 7.01 4 2.64 3.63 3.79 6.85 1.28 4 2.83 4.18 2.56 3.38 4.80 4 2.34 3.33 3.82 2.44 4.10 4 2.34 3.33 3.82 2.44 4.10 4 4.67 5.81 2.55 4.74 3.85 4 4.67 5.81 2.55 4.74 3.85 4 4.67 5.81 3.36 5.17 4.30 4 1.73 6.29 3.11 1.20 3.29 4 1.21 4.61 5.01 4.80 4.57 4 5.81 3.02 3.15 2.19 2.78 5 4 2.09 3.77 3.86 1.82 5.82 4 2.09 3.77 3.84 1.82 5.82	AAB39275 brain N/A 0.61	brain N/A 0.61	N/A 0.61	0.61		*	22	*	2.08	000) ;		3	3.71	•
4 5.19 5.14 5.21 5.14 5.21 4 2.57 4.47 5.19 5.29 7.01 4 2.93 4.18 2.58 3.38 4.80 4 2.34 3.33 3.82 2.44 4.10 4 2.34 3.33 3.82 2.44 4.10 4 2.34 3.33 3.82 2.44 4.10 4 4.67 5.81 2.55 4.74 3.85 4 4.67 5.15 3.36 5.17 4.30 4 4.16 2.79 3.62 5.06 3.32 4 1.73 6.29 3.11 1.20 3.29 4 1.21 4.61 5.01 4.80 4.57 4 5.81 3.02 3.15 2.19 2.78 5 5 3.77 3.86 1.82 5.82 3 3.77 3.84 1.82 5.82	R76247 brain 0.43	brain 0.43	0.43	0.43		÷	×	•	9.05	3.34	¥ 8	<u> </u>	86.6	28 28	4
4 2.57 4.47 3.19 6.85 1.28 4 2.93 4.18 2.58 3.38 4.60 4 2.94 4.18 2.58 3.38 4.60 4 2.34 3.33 3.82 2.44 4.10 4 4.67 5.61 2.55 4.74 3.85 4 2.92 5.15 3.36 5.17 4.30 4 4.16 2.79 3.62 5.06 3.32 4 1.73 6.29 3.11 1.20 3.29 4 1.21 4.61 5.01 4.80 4.57 4 5.81 3.02 3.16 2.19 2.78 5 3.77 3.86 1.92 5.82		1.23	8.			₹	28	→	5.19	4 .	5	2 6	7.01	0.18	*
4 640 3.53 3.54 3.55 3.58 4.60 4 2.93 4.18 2.58 3.38 4.60 4 4.67 5.61 2.55 4.74 3.85 4 2.82 5.15 3.36 5.17 4.30 4 2.82 5.47 3.81 5.36 5.30 4 1.73 6.29 3.11 1.20 3.29 4 1.21 4.61 5.01 4.80 4.57 4 5.81 3.02 3.15 2.19 2.78 5 3.77 3.86 1.82 5.82	0.05	t-cell 0.05	90.0			0.	•	→	2.57	18.4		28.8	1.28	0.83	*
4 2.83 3.18 2.50 4 2.34 3.33 3.82 2.44 4.10 4 4.87 5.81 2.55 4.74 3.85 4 2.82 5.15 3.36 5.17 4.30 4 4.16 2.79 3.62 5.06 3.32 4 1.73 6.29 3.11 1.20 3.29 4 1.21 4.61 5.01 4.80 4.57 4 5.81 3.02 3.15 2.19 2.78 5 3.77 3.86 1.92 5.82 3 3.77 3.84 1.82 5.82		0.28				2	_	*	6.40	3.5	. c	88.6	9.4	4.50	•
4 4.67 5.61 2.55 4.74 3.85 4 2.82 5.15 3.36 5.17 4.30 4 4.16 2.79 3.62 5.06 3.32 4 1.73 6.29 3.11 1.20 3.29 4 1.21 4.61 5.01 4.80 4.57 4 5.81 3.02 3.16 2.19 2.78 3 3.02 3.77 3.86 1.82 5.82	N25920 - 1.81	. 1.81	1.81			ĕ	92	❖ ·	2.83	<u> </u>	26	244	4.10	5.95	•
4 4.67 5.81 2.30 5.17 4.30 4 2.82 5.15 3.36 5.17 4.30 4 4.16 2.79 3.62 5.06 3.32 4 1.73 6.29 3.11 1.20 3.29 4 1.21 4.61 5.01 4.80 4.57 4 5.81 3.02 3.16 2.19 2.78 4 2.09 3.77 3.86 1.82 5.82		brain 0.93	0.93			8	~	•	K.3	3 1	200	7.7	3.85	0.49	*
4 2.02 5.15 5.36 5.17 5.34 5.17 5.36 5.37 5.37 5.39 5.47 5.41 5.39 5.47 5.39 5.47 5.39 5.47 5.41 5.49 5.45 5.41 5.01 4.80 4.57 5.49 5.81 5.02 5.49 5.78 5.82 5.82 5.82 5.82 5.82 5.82 5.82 5.8	3.58	3.58	3.56			13.0	_	*	4.67	5.61	8 6	, , , , , , , , , , , , , , , , , , ,	Ş 7	8.	*
4 4.16 2.79 3.62 6.06 3.32 4 3.55 5.47 3.81 5.38 2.39 4 1.73 6.29 3.11 1.20 3.29 4 1.21 4.61 5.01 4.80 4.57 4 5.81 3.02 3.16 2.19 2.78 4 2.09 3.77 3.86 1.82 5.82	AAR30104 bone Y 0.98	bone Y 0.96	96:0 ×			3.5	_	₹	2.82	5.15	3.36	7.6	8.	5	. •
4 3.55 5.47 3.81 5.38 2.39 4 1.73 6.29 3.11 1.20 3.29 4 1.21 4.61 5.01 4.80 4.57 4 5.81 3.02 3.16 2.19 2.78 4 2.09 3.77 3.86 1.82 5.82 3	28.0	0.83				3.03		•	4.16	2.79	3.62	90.0	3.32	28.7	•
4 1.73 6.29 3.11 1.20 3.29 4 1.21 4.61 5.01 4.80 4.57 4 5.81 3.02 3.16 2.19 2.78 4 2.09 3.77 3.86 1.82 5.82	10.52	10.62	10.52	•	•	38.0	9	*	3.55	5.47	3.81	5.38	2.30	1.1	• •
4 1.21 4.61 5.01 4.80 4.57 4 5.81 3.02 3.16 2.19 2.78 4 2.00 3.77 3.86 1.92 5.82	SOC CONTRACTOR CONTRAC	8000	800			•	3	4	1.73	6.29	3.11	1.20	3.29	80 ·	•
4 5.81 3.02 3.15 2.19 2.78 4 2.09 3.77 3.84 1.92 5.82	ANASTON INSURAL STATE OF THE ST	Netical Asset 14.33	****				82	•	1.21	4.61	5.01	4.80	4.57	1.28	•
4 2.09 3.77 3.86 1.92 5.82 3	SOUTH CAN BOOK DOWN THE STATE OF THE STATE O			1 5		_	45	4	5.81	3.02	3.15	2.19	2.78	1.53	•
		96'0 V/V	96°0	8 6			. 8	- →	2.09	3.77	3.86	1.92	5.82	3.66	•
									60						

Table 2-2: T3NO vs. BPH: Gene upregulated 5 701d or higher in at least 2 tumor samples, or 3 fold or higher in at least 3 tumor Samples.

	IMAGE	Gen Bank	Tissue	Secreted/Me		,	Fold(AveTumoria Fold(MPM120/A Fold(MPM121/A Fold(MPM124/A Fold(MPM122/A Fold(MPM123/A Fold(MPM125/A # of Tumors	Fold(MPM120/A	Fold(MPM121/A	Fold(MPM124/A	Fold(MPM122/A	Fold(MPM123/A	Fold(MPM125/A	t of Tumors
ğ Ö	Clone 1D	Accession	Distribution	mbrane Bound	Avenda	AVe I umor	veBPH)	veBPH)	VeBPH)	VeBPH)	(week)			
Ş	******	******		4774	88	10.39	•	3.82	5.10	3.87	3.79	2.79	1.62	•
3		A4404800	•	Š		42.4	· e.	5.39	30.0	3.53	2.17	2.42	4.40	- *
124	281377	N47886	Bone	;	8 3		· ~	244	3.12	4.85	57.7	3,48	6.29	*
125	383823	AA704650		>	5 6	9:15	າ ແ	2.28	5.31	3.00	4.08	4.15	2.08	•
126	395400	AA757401	pove	;	07.7	8 8	, c	7.1	4.81	3.90	3.23	4.16	2.65	•
127	138507	R63253	•	> :	Ę.	8 6	, «	3.37	90.50	1.38	1.50	60.0	3.28	*
128	289071	N63609	,	> [9.91	8.5 5.5		181	4.23	4.13	4.28	4.83	1.56	•
1 20	460203	AA877461	•	×	6.5	3 0.0	, c	5.12	6.73	3.43	3.25	0.92	0.85	•
8	242061	H80333	testis		. ;	7.6		302	4.77	30.00	2.28	4.65	2.52	•
Ē	207098	H48502	empolo		0.10	2 2	, ,	1.87	5.82	3.24	4.60	39.	3.25	*
<u> </u>	396839	AA758154	colon		9 1 5	57.7	, ,	70.	3.28	4.07	1.88	4.90	3.98	4
ડ	208381	H64095	colon	>	0.97	07.5	, c	5.5	3.91	5.86	3.28	1.47	3.16	•
ž	23897	R38364	•		80 ·	* 1	, ,	187	3.82	\$35.	3.37	3.97	2.04	+
135	1467905	AA883504	thymus	₹	÷.	# H	n (94.6		3.20	06.0	4.48	1.12	*
5	1556526	AA835560	prostate	~	1.17	3.77	77	00.0	8 6		85	3.33	3.88	*
137	401418	AA115537	breast		9.5	1.72	က	1.82	3.32	0 ti	8° 7		1.80	•
\$	744001	AA629039	heart	¥	3.87	12.24	ဗ	1.80	3.05	3.2/	D	- 1	7	
130	283110	N63848	•		3.38	10.70	n	2.38	đ.	4; 4;	3.86	8 6	5. 6	• •
3	782047	AA429802	prostate	Maybe	1.17	3.67	60	3.26	2.51	3.73	2.89	3.22	17.6	• •
3	454880	44677208			1.02	3.16	6	2.09	3.15	1.60	3.87	3.10	4.54	•
: 5	184203	44700370		* // *	0.50	53	n	4.26	3.84	4.06	3.20	5 .	1.87	•
<u>:</u> ;	7076	2777	·	<u> </u>	8	5	ю	2.04	36.6	3.79	3.14	3.71	<u>5</u>	•
2 :		70000			3 8	2	· 67	3.33	3.10	2.05	4.16	0.88	4.58	•
	70534	***************************************	esensord.	;	3 .	3	· e	2.11	2.81	3.08	3.17	3.53	3.40	*
5	126/85	R10015	F .	-	<u>.</u>) e	3.22	3.29	1.89	3.35	277	4.5	~
5	420200	AA007370	Dreast 1		9 5	6.67) r	3.15	2.85	3.25	3.12	1.85	3.33	•
4	136819	R36465	•		Y) 97 467		3,66	5.22	3.12	123	0.47	3.55	•
2	433350	AA700604			42.83	91.831	? (2 6	463	100	1.81	0.36	3.03	4
2	755276	AA496349	brain			0.30	n	70.0		£ .	16.41	13.45	1.13	•
3	270017	N32514			0.0	0.23	92 :	1 5	2 5		0.61	3.63	22.02	60
15	782233	AA431721	prostate	>	0.20	3.10	15	24.92	4 1	: t	95. 55	3.78	2.68	n
152	320495	W16659	pone		90.0	98.0	‡	1.39	3.75	78.0	0°-1	2077	2.05	• •7
53	203579	N94111	spieen		98.0	12.90	13	0.73	1.52	77.4	14:07			
2	281039	717717	endotheliei		0.78	8	13	43,16	25.22	5.01	222.	S	7 6	, ,
155	490718	AA115761	empvo		0.0	0.46	12	46.34	6.67	. 83	0.27	18.31		, (
35	796747	AA443147	eloq		90.0	0.92	F	0.12	0.12	2.23	\$5.00	5.81	\$ 100	, (
5	287728	NSD158	•	>	0.51	5.43	£	20.89	31.93	7.51	1.25	9.86	1.67	n (
\$	282080	N48259	etetecnu	•	8.	10.34	5	21.89	29.40	4.51	1.88	1.07	2.28	n (
3 \$	7,4700	44446627	,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,		80.7	33.97	80	14.84	26.99	1.81	1.28	0.78	† 71	n
3 5	470218	4403997	nmetate		2 05	23.62	60	1.97	4.75	30,1	5.39	32.26	2.68	n
3 ;	210246	100000			200	0.67	sc	3.54	3.42	1.83.	1.38	34.08	2,42	n
ē ;	7907	Logon .			5	3	, ,-	0.87	3.83	1.15	32.30	5.68	0.12	n
ž !		1/0840		;	? 5	4.5		4.67	30.78	1.10	3.88	0.00	1.38	60
3	19/43	A43/241	•	Š	Ì		. ,	376	7.30	2.60	21.59	8.58	1.59	n
ž	486035	AA040861	made	Š	5.25	35.88	~		•	i				
								_						

Table 2-2: T3NO vs. BPH: Gene upregulated 5 fold or higher in at least 3 tumor Samples.

Distribution Bound 0.08 0.48 0.04 0.23 0.04 0.23 0.04 0.23 0.04 0.23 0.04 0.05 0.04 0.23 0.04 0.05 0.04 0.05 0.04 0.05 0.04 0.05 0.04 0.05 0.04 0.05 0.04 0.05			Gen Bank	į	Secreted/Me			Fold(AveTumor/A	Fold(MPM120/A	Fold(MPM121/A	Fold(MPM124/A	veBPh)	Fold(AveTumor/A Fold(MPM120/A Fold(MPM121/A Fold(MPM120/A	VeBPH)	Samples
		MAGE	Accession	Distribution	mbrane	AveBPH		VeBPH)	(MagAu)	î i		84	1,47	3.24	n
ACCISION COMMUNICATION COMMUNICATIO		5000	Number		Bound	Ş	87.0	60	13.15	16.83	1.72	9. °C	2.26	0.81	n
Available align Old		298417	N74131	coloa		90.0	8.5	•	6	7.96	4,25	20.28		27.68	n
Wilson Visitaria 0.02 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.02 0.01 0.02 0.01 0.02 0.01 0.02 0.01 0.02 0.01 0.02 0.01 0.02 0.02 0.02 0.03		842718	AA233809	skin		9.0	0.23	ρ (, t	4,23	0.65	4.75	96.0	7.5	•
WESSER CORRESPONDED 3.32 0 1.15 1.25 1.05 2.25 1.05 2.25 1.05 2.25 1.05 2.25 1.05 2.25 1.05 2.25 1.05 2.25 1.05 2.25 1.05 2.25 1.05 2.25 1.05 2.25	-			,		0.02	0.1	Ð	2 ;	4	2.16	16.65	90.	\$	• •
March Marc		10/10/	AMS/ 192	•		0.59	3.32	9	V.1	3 4	177	1.32	30.	2.26	3 (
Accessing problemations SERIES SERIES CORRES		20108 80108	R92362	• 1		180	4.58	80	5.88	80.8L		0.72	0.88	. 2.12	n
Accessing some of the control of the contro		687667	AA235370	hypothalamus		5 5	32.98	80	6.81	14.24	9.07	0.13	17.97	0.10	e
W1647678 Doors CAST STATE S		712950	AA282273	₩ •		2 8	570	so	6.26	0.51	7/1	5	0.89	2.62	n
Holista command Color 1,17 6,00 1,27 2,12 2,13 2,10 1,10 6,10 1,10 1,10 1,10 1,10 1,10		504761	AA148735	pou		3 6	÷ +	167	3.13	4.09	0.73	30,4	11.48	6.86	n
MATCHEST LOSE 3231 2.22 2.25 8.25 0.00 1.75 8.25 9.25 1.75 8.25 9.25		322443	W16424	adrenal		22	57.	, u	2.72	4.85	2.60	9		828	•
Averages (state) tests (172 8.6 7.26 1.26 1.17 0.00 1.13 2.88 1.26 1.26 1.27 1.19 0.00 1.13 2.88 1.26 1.26 1.19 1.19 1.19 2.88 3.48 0.00 1.13 2.88 3.48 0.00 1.13 2.88 3.48 0.00 1.13 1.19 2.88 3.24 0.00 1.13 2.88 3.44 0.00 1.13 2.88 3.44 0.00 1.13 2.88 0.00 1.13 0.00 1.13 4.48 1.13 4.48 1.13 4.48 1.13 4.48 1.13 4.48 1.13 4.48 1.13 4.48 1.13 4.48 1.13 4.48 1.13 4.48 1.13 4.48 1.13 4.48 1.13 4.48 1.13 4.48 1.13 4.48 1.13 4.48 1.13 4.48 1.13 4.48 1.13 4.48 1.13 4.44 4.44		9696	0.46448	•		0.0	3.31	n		2.75	8.08	0.00	7.48		
Accordant Mary Mary College Date of Mary Mary College 148 734 6 128		200	450110			1.72	8.60	ເດ	7.52	;	2.17	0.90	1.13	48.6	, ,
Avidence bone 0.71 47.00 6 1.82 1.40 1.50 0.00 7.22 4.38 9.35 Avidence one 0.71 47.00 6 1.80 1.42 1.43 1.43 0.00 7.22 4.38 1.43 1.44 1.43 1.44 1.45 1.44 1.45 <th< td=""><td></td><td>782537</td><td>AA448484</td><td>1000</td><td>></td><td>87</td><td>7.3</td><td>10</td><td>7.68</td><td>21.21</td><td>: 3</td><td>12.13</td><td>2.89</td><td>3.52</td><td>7</td></th<>		782537	AA448484	1000	>	87	7.3	10	7.68	21.21	: 3	12.13	2.89	3.52	7
AMAZESSES with pasts with pas		277596	N57002	euoq	-	2	47.00	10	1.82	8.	1	8	7.22	4.30	n
Mailant		773208	AA425850	skin			9	16	1,52	1.24	13.33	3	4	0.89	n
Highest High		364865	AA035745	breast		\$ 1	90.4		1.96	14.29	6 .	8	*	2.65	n
		247587	N58136	bone		0.01	0 0 0	, ·	69	.00	6.74	3.	3 3	5	63
National			0.007	•		0.14	0.67	o		72.44	4.47	1.37	3	2 9	
AUSTAGES Dome 11274 SEGD 5 147 2.55 8.45 8.14 0.24 0.28 2.34 MST4585 Domin 0.02 0.02 1.47 8.45 8.19 0.64 0.28 2.34 MST4585 Demin 0.02 0.02 1.72 8.45 8.19 0.64 0.28 2.34 MST4585 Demin 0.03 0.10 1.72 8.45 8.19 0.64 0.28 2.34 0.39 1.87 0.39 0.34 1.30 0.39 0.34 1.30 0.39 0.34 1.30 0.34 0.34 1.30 0.39 0.34 1.30 0.39 0.34 1.30 0.39 0.34 1.30 0.34 1.30 0.34 1.30 0.39 0.34 1.30 0.34 1.30 0.34 0.39 0.34 0.34 0.34 0.34 0.34 0.34 0.34 0.34 0.34 0.34 0.34 0.34 0.34		8	200	atracata		90.0	0.36	ın	79.7	1 40	5.37	13.00	2.45	3.36	, (
AMSTASS Dominal Paris O.02 5 1.87 6.45 0.54 0.23 2.34 A.54 0.54 0.24 0.39 2.34 A.54 0.64 0.24 0.39 2.34 A.54 0.64 0.24 0.39 0.34 A.54 0.64 0.64 0.24 0.39 0.34 4.45 0.69 0.54 0.74 0.64 0.74 0.64 0.74 0.64 0.74 0.64 0.74 0.64 0.74 0.64 0.74 0.64 0.74 0.64 0.74 0.64 0.74 0.64 0.74 0.64 0.74 0.64 0.74 0.64 0.74 0.64 0.74 0.64 0.74		72365	20101			12.74	58.08	vo.	1,47	.	2 65	8.18	1.05	7.68	•
MATORIAS briant 7.34 53.36 5 7.28 8.45 5.17 6.53 2.43 0.38 HISADORIA briant 7.34 53.36 5 7.28 8.45 7.29 6.53 2.43 0.39 1.53 0.34 1.13 0.34 1.13 0.34 1.13		113204	AA425692	•uoq		6	000	٠,	1.67	6.21	3 9	380	0.26	2.36	n
AAAGOAGAS Drain AAAGOAGAS Drain AAAGOAGAS Drain AAAGOAGAS Drain AAAGOAGAS Drain AAAGOAGAS Drain AAAGOAGAS AAAGOAGAS AAAGOAGAS AAAGOAGAS AAAAGOAGAS AAAAGOAGAS AAAAGOAGAS AAAAGOAGAS AAAAAGAAGAA AAAAAGAAGAAAAAAAAAAAAAAAAAAAAAAAAAAAA		277327	N57483	ciero			33.36	ĸ	7.28	8.45	3 1		2.43	0.38	n
H1944 H1954 H1956 H195 H		770066	AA430645	brain		8	3 5	•	1.90	8.33	7.29	3 3	3	1.83	n
N/2659		51620	H19446	•		9.0	0.12	• •	8.62	9.86	4.53	4 0		1.27	n
W02428		266343	N26556	•		0.70	3.70	• •	4.09	12.69	6.34	9.0		8.83	es
Mo6139 Drain 0.05 0.22 4 0.05 0.22 4 10.35 1.67 0.08 8.74 0.07 4 AA/12407 breast 0.02 0.07 4 10.35 0.16 0.28 2.13 8.96 4.11 NG44570 bing 0.09 0.22 4 0.046 1.09 0.28 2.13 1.51 1.74 3.80 4.11 NG44570 bing 0.03 0.22 4 0.046 1.09 0.05 0.17 1.76 0.06 0.07 1.00 0.04 1.00 0.04 <td></td> <td>296170</td> <td>W02426</td> <td>•</td> <td></td> <td>5</td> <td>7.8/</td> <td>• •</td> <td>0.98</td> <td>7,96</td> <td>1.80</td> <td>5.23</td> <td>1 2</td> <td>2.67</td> <td>n</td>		296170	W02426	•		5	7.8/	• •	0.98	7,96	1.80	5.23	1 2	2.67	n
AA728407 breast 0.02 0.07 4 10.35 0.16 0.28 2.13 0.09 2.13 0.09 2.13 0.09 2.13 0.09 2.13 0.09 0.28 4 10.35 0.16 0.28 2.13 1.01 1.74 <		278504	N66139	brain		9	27.0	, ,	330	1.67	96'0	9.74	3 8	157	n
AAA45470 lung 0.06 0.28 4 0.46 1.96 0.68 3.72 15.21 15.21 15.21 15.21 15.21 15.21 15.21 15.21 15.21 15.21 15.41 17.4		K28587	AA128407	breast		0.02	0.07	•	£0.35	0.16	0.26	2.13	S :	: C	- 11
Noestee 0.05 0.22 4 7.22 8.76 3.67 2.18 1.01 1.14 N73563 0.042 1.81 4 7.22 8.76 5.57 1.28 1.09 6.79 N73563 1.eath 0.61 2.18 4 7.69 2.40 6.57 1.28 1.92 6.79 H75655 postate 0.61 2.18 4 7.69 2.40 6.57 1.28 1.92 6.79 A4463737 breath 4.06 2.042 4 3.45 0.81 2.40 2.24 6.79 1.39 A4463737 breath NA 2.86 11.62 4 0.28 1.40 2.24 5.89 1.33 AA46736 brone V 1.06 4.28 4 0.28 2.40 6.78 2.44 6.79 1.39 4.70 1.33 1.13 2.44 6.79 2.24 6.79 1.39 1.13 2.44 6.79		800526	AA454570	auni		90.0	0.28	•	970	8	0.68	3.72	15.01	3	· et
Harrist Harr		#y0000	N80400	•		0.05	0.22	*	2 6	9.78	3.67	2.18	1.01		, e.
H37555 prosition 0.51 2.18 4 7.99 7.90 0.66 0.00 0.68 8.75 7.50 7.50 7.50 7.50 7.50 7.50 7.50 7		20K007	N73583	•		0.42	1.81	*	7 7	2.40	5.57	1.28	1.92	9.0	. "
T49325 Testis 0.01 0.06 4 3.45 0.81 2.88 7.56 1.37 882 AAA48373 breast 4.96 20.42 4 3.45 0.81 2.88 7.47 8.01 2.44 1.39 AAA48373 breast 4.96 20.42 4 2.09 6.18 6.48 2.24 5.89 1.35 AAA6736 iung Y 1.06 4.28 4 10.98 2.50 4.10 3.36 1.15 2.06 AAA67365 iung Y 1.06 4.28 4 10.98 2.50 4.10 3.36 1.15 2.06 AAA676604 bone 0.04 0.14 4 2.52 2.29 7.11 2.63 4.00 AAA676604 bone 0.03 0.13 4 2.24 1.36 2.24 4.00 AAA6302 heart 0.04 1.79 4 2.62 2.29 7.11 2.63			MEYERS	atatata		0.51	2.18	→	80. V	7 00	0.68	0.80	99.0		
AAA65373 breast 4.96 20,42 4 3.45 0.01 7.47 8.01 2.44 1.39 AAA65373 breast 1.62 4 3.07 1.98 7.47 8.01 2.44 1.39 AAA63736 broads 1.58 6.38 4 2.09 6.18 6.46 2.24 6.59 1.52 AAA67365 bring Y 1.06 4.28 4 1.08 2.50 4.10 3.36 1.15 2.06 AAA67336 bring Y 1.06 4.28 4 2.22 7.11 2.63 4.39 4.50 AAA67634 bring Q.04 0.14 4 2.52 2.29 7.11 2.63 4.99 4.00 AAA63735 brind Q.04 0.17 4 2.93 7.71 2.63 4.99 4.00 AAA63735 brind Q.03 Q.10 4 Q.24 1.86 1.30 4.13		070007	2007			0.0	90.0	₹	A7.C		2.68	7.56	1.37	8.82	, ,
AMMS373 United 11.62 4 3.07 130 6.18 6.46 2.24 5.69 1.52 AMM37365 Iung Y 1.06 4.28 4 2.09 6.18 6.46 2.24 5.69 1.15 2.06 AMM37365 Iung Y 1.06 4.28 4 1.038 2.50 4.01 3.36 1.15 2.06 AMM37365 Iung Y 1.06 4.28 4 2.62 2.29 7.11 2.63 4.00 4.00 AMM37360 Iung Y 1.06 0.25 4 2.62 2.29 7.11 2.63 4.00 AMM37375 Iung Iung 1.79 4 2.62 2.29 7.71 2.63 4.00 AMM37375 Inset Iung Iung 4 3.05 1.36 1.36 1.30 1.30 AMM37375 Iung Iung Iung Iung Iung Iung		196/9	148323	97.00		8.	20.42	*	3,45	. o.	17.0	8.01	2.44	4.30	n (
AA101878 prostate V 1.58 6.38 4 2.09 6.18 0.00 4.10 3.36 1.15 2.06 AA487365 Ling Y 1.68 4.28 4 10.38 2.50 4.10 3.36 1.15 2.03 4.00 AA487365 Ling Y 1.06 0.14 4 0.28 7.63 0.28 7.11 2.63 4.00 AA4873604 bone 0.04 0.14 4 2.52 2.29 7.11 2.63 4.00 AA483735 Ling 1.78 4 2.83 3.44 6.26 7.90 1.68 2.22 AA483730 heart 0.03 0.10 4 3.05 1.55 3.64 1.86 1.30 4.10 AA48302 bone 0.01 0.05 4 0.44 0.74 0.74 0.74 0.74 AA47522 cervix 0.36 0.31 4 0.36 0.36		811028	AA485373		4/14	2.86	11.62	*	3.07	9.	. 49	2.24	5.69	1.52	n ·
N64786 VAMSTAGE <		489637	AA101878	prosiding	2	2	6.38	4	5.09	6.18	P	25	1.15	2.06	n
AAA87365 Ling 7,00 0.14 4 0.28 7,63 0.28 4.99 4.00 AAA73064 bone 0.04 0.14 4 2.52 2.29 7,11 2.63 4.99 4.00 AAA83735 Lost 0.04 0.179 4 2.93 3.44 6.26 7.90 1.68 2.22 AAA3300 heart 0.05 0.10 4 3.05 1.55 3.64 1.88 1.30 0.74 0.74 AAA3300 heart 0.03 0.10 4 3.91 7.78 1.46 8.70 0.74 0.74 AAA30620 AAA4300 heart 0.05 4 6.41 6.16 0.96 0.36 0.34 AAA47522 carvix 0.36 1.39 4 8.55 7.01 2.57 0.06 0.88 3.47 AAA130857 Neuron N/A 1.17 4.49 4 2.76 2.15 2.15 2.1		284624	N64796		;	3 5	4.28	4	10.98	2.50	01.4	3 5	0.79	5.34	n
AA676604 bone U.O.S. C.T.T. 2.52 2.29 7.11 2.50 1.68 NA7012 NUA 0.06 0.25 4 2.63 3.44 6.26 7.90 1.68 AA83735 heart 0.03 0.10 4 3.05 1.55 3.64 1.89 1.83 AA630620 heart 0.01 0.05 4 3.91 7.78 1.46 8.70 0.74 AA630620 AA630620 0.01 0.05 0.11 4 6.41 6.16 0.96 0.35 1.33 AAA47522 cervix 0.36 1.39 4 1.91 3.10 4.57 10.12 1.60 AA130857 Neuron NUA 1.17 4.49 4 2.76 2.15 2.16 8.17 4.00		841507	AA487385		-	3 3	3	•	0.28	7.63	0.28	3	8	8;4	n
N47012 NUA 0.06 0.25 4 2.93 3.44 6.26 7.30 7.31 7.30 7		882497	AA676604			5 6	, u	4	2.52	2.29	7.11	3 3	5	222	n
AAA83735 N/A 0.46 1.78 1.65 3.64 1.86 1.25 AAA43300 heart 0.03 0.10 4 3.91 7.78 1.46 8.70 0.74 AAA30620 AAA43062 0.03 0.11 4 6.41 6.46 0.96 0.35 1.39 AAA47522 cervix 0.36 1.39 4 8.55 7.01 2.57 0.65 0.88 AAA130857 bone 0.36 1.77 4.49 4 2.76 2.15 2.16 8.17 4.00 AA130857 Neuron N/A 1.17 4.49 4 2.76 2.15 2.15 8.17 4.00		280155				8	C. 7.	• •	2.93	3.44	6.26	8.	3 5	11.30	n
AAA43300 heert 0.03 0.10 4 3.91 7.78 1.46 8.70 0.14 AA29283 bone 0.01 0.05 4 6.41 6.16 0.96 0.35 1.33 AAA30820 AAA4752 cen/x 0.36 1.39 4 8.55 7.01 2.57 0.65 0.88 AAA4752 cen/x 0.36 1.39 4 2.76 2.15 8.17 4.00 AA130857 Neuron N/A 1.17 4.49 4 2.76 2.15 8.17 4.00		1466788	Ī		Y	0.46	8/:	•	3.05	1.55	3.64	38.		0.74	n
AA530626 bone 0.01 0.05 1.33 AA530620 0.03 0.11 4 8.15 7.01 2.57 0.65 0.88 AA447522 cervix 0.36 1.39 4 1.91 3.10 4.57 10.12 1.60 AA447524 cervix 0.36 1.39 4 2.76 2.15 2.16 8.17 4.00 AA130857 Neuron N/A 1.17 4.49 4 2.76 2.15 2.16 8.17 4.00		784589	-			0.03	0.10	•	3.91	7.78	1.48	8.70	• ()	, C	n
AA630620 AA630620 AA447522 cervix 0.36 1.39 4 8.55 7.01 2.57 0.65 0.88 AA447522 cervix AA001870 bone AA130857 Neuron NJA 1.17 4.49 4 2.76 2.15 8.17 4.00		772077				0.0	0.05	•	44	6.16	96.0	0.35	E	2 5	
AAA47522 cervix 0.36 1.39 4 8.55 7.01 4.57 10.12 1.60 4.00 4.00 4.57 10.12 1.60 4.00 4.00 6.31 6.31 6.31 6.00 6.00 6.00 6.00 6.00 6.00 6.00 6.0						800	0.11	•	÷ (2.57	0.65	0.88	3.47	, (
AAA47522 centx 2.28 8.77 4 1.91 3.10 4.00 AA001870 bone 2.28 8.77 4 2.76 2.15 8.17 4.00 AA130857 Neuron NJA 1.17 4.49 4 5		11000				0.36	1.39	4	8.55	5 6	197	10.12	1.60	1.76	n (
AA001870 bone		782575				2.28	8.77	4	1.94	0.10		8.17	0.4	3.75	n
AA130857 Neuron NJA 1.17		428166	-		į		449	*	2.76	2.15	2.10	;			
Sign Company of the C		586826	_		₹	<u> </u>	ř		,						
									જ						

Table 2-2: T3NO vs. BPH: Gene upregulated 5 fold or higher in at least 3 tumor samples or 3 fold or higher in at least 3 tumor Samples.

				Secreted/Me			4	A)001400VA	Fold/MPM121/A	Fold(MPM124/A	Fold(MPM122/A	Fold(MPM123/A	Fold(MPM125/A	# of Tumors
Order	Clone 1D	•	Tissue Distribution		AveBPH	AveTumor	Fold (Ave Lumoria Folding veBPH) veBPH) veBPH) VeBPH) Samples	veBPH)	veBPH)	veBPH)	veBPh)	veBPH)	veBPH)	Samples
				Bound	6	6	•	7.50	0.49	0.49	7.80	2.20	4.30	6
204	730838	AA417012	testis		0.02	80.0	•	S: 6		1 10	2.48	8.86	2.62	- m
8	837953	AA458578	Bunj	>	28.	29.78	•	3.70	28.0			5.74	6.57	6
200	503824	AA133959	orain		0.12	9.4	*	1.62	0.0	0 1	9 0	3.0	2	es
2,0	208210	H85286	placenta		0.89	3.34	*	5.28	5.92	7.33	1.6¥	2 1		
;	427006	AA001841			0.11	0.43	•	3.08	.05 20.	3.0		16.0	3 5	
; ;	******	700,000	1		184	06.90	4	0,50	0.29	0.34	4.24	12.61	00.4	, (
717	8/3	4010K	The Carlot	****	2 2	52.21	•	1.02	1.89	3.14	8.23	5.08	2.80	n (
213	3 5	W/0348		Ç	3	-	•	2.83	6 .04	2.57	2.67	3.32	4.58	n :
27.4	143062	K71336			ţ •	5 6		5.05	2.39	2.70	1.07	4.57	6.14	n
215	131588	R23727	•	:	2 6	3 9	• •	2.83	300	1.92	4.91	6.33	5.22	m
218	240151	H82706	poue	Maybe	3 :	*	• •	3.4	7.49	3.46	2.29	3.62	2.52	n
217	452116	AA707219	pone		1	5 6	• -	, e	9	2.14	6.14	2.91	2.55	n
218	201125	R98472	skin	₹	3	E :	• •	9 9	4.18	2.20	4.20	1.20	2.78	n
218	126713	R07115	•		0.27	86.0	•	Š	2 8	1.69	2.78	7.17	2.61	
82	768179	AA424849	brain	4	52	80.0e	•	\$ 6	9 9		3.7	2.33	2.51	n
ជ	250678	H95976	gun	Maybe	6 .0	14.42	₹	2.85	000	57.5		157	1.47	n
Ħ	273499	N33263	•		0.0	0.32	4	3.68	2.67	6.75	6	ā	1.82	n
S	755765	AA496455	muscle		1.95	6.84	*	2,12	3.80	0 ti		Š	7.18	•
Ř	815047	AA465166	•uoq		2.68	9.34	es	2.62	9.	28.1	7.4	5 6	2.03	. 63
225	384670	AA709333		**	4.17	14.53	6	3,11	2.41	3.77	8 7	. .	2 5	, es
228	782200	AA431964	muscle		1.53	5.30	က	2.48	6.32	2.54	1	07.7		•
222	82730	M29500	hrein		3.0	22.92	က	2.33	2.11	4.07	6.25	3.48	# · · · · ·	
į	126311	805810	9	X	1.92	6.62	ဗ	1.07	2.84	3.69	89.68		8 1	, ,
3 8	770880	20772744	Jecombe		80.8	27.68	n	6.28	3.15	7.87	1.10	0.49); i	, ,
3 8	2000	20000			8	8	n	1.43	3.68	2.58	5.27	5.89	1.71	D (
8	90706/	/200007			3 6	707	· 67	2.91	3.75	7.21	4.16	1.12	4.30	n
ā i	9	Hooks	Dioseisie.	2	2 6	2	, es	9.82	3.38	1.53	3.96	0.90	08:0	n
22	611373	AA176833	oun.		5 6	\$ 8) e	1.86	1.13	5.78	5.41	1.13	5.00	•
ន្ត	325380	AA284285	spies.		5 6	3 6	, r	5.38	1.97	4.32	1.12	1.56	6.01	60
ផ	434822	A703114	•		3 1	7.77		80 7	4.75.	2.95	4.56	1.19	ă.	n
22	1049033	AA778675	7 .		1.87	9 9	, c	2 7	88.	285	5.61	3.78	1.72	•
ន្តី	882483	AA676598			0.70	3 5	י כ	5 6	0.74	1.14	3.10	10.17	4.12	•
ដ	767130	AA424535	kidney		340	10.1	· •		77	373	14.1	0.36	221	n
2	126321	R06479			8 1	2 6	.	7.07	78.6	4	2.89	8.21	0.15	•
ğ	23046	H76328	•		0:0	0.22	? (. S	8 6	2.87	5.02	2.83	1.90	•
3	624271	AA181179	Poue	\$	6.78	22.81	, c	2.5	0 13	7 7 7	0.68	96.0	0.68	n
241	123265	R00284	lymph	₹	5.30	7.72	, c	77.4	2 6		5	3.53	5.10	n
242	367236	W\$3638	٠		0.72	2,40	60	5.76	2.05	2 9	<u> </u>	3.48	2.80	n
243	123331	R00479	prostate		1.03	3,43	က	2.37	4.72	8 .			121	•
7	240694	H78134	,	>	1,85	6.17	က	2.79	3.07	1.59	S :	0.50	1 0	• •
245	347182	W80611	brain		0.02	0.07	ဗ	0.45	5.03		*	7.0	,	, ,
248	62002	H23232	postate		89	0.11	ဗ	1 .	4.38	6.28	4.13	79.2	; ;	•
247	160628	H25019	domy	>	0.41	5.7	ო	2.02	4.53	3.17	4.45	2.72	2	, ,
	1505294	AAGOERIA	teaffs	N/A	13	3.66	ဗ	1.98	2.43	7.21	1.55	3.28	X	,
ł					:			œ						

Table 2-2: T3NO vs. BPH: Gene upregulated 5 told of higher in all east 2 tumor samples or 3 fold or higher in at least 3 tumor Samples.

			i	Secreted/Me	,		Fold(AveTumor/A	Fold(AveTumor/A Fold(MPM120/A Fold(MPM121/A Fold(MPM122/A Fold(MPM122/A Fold(MPM122/A Fold(MPM120/A	Fold(MPM121/A	Fold(MPM124/A	Fold(MPM122/A veBPh)	Fold(MPM123/A	veBPH)	Samples
ode.	MAGE	Accession	Distribution	mbrane	AVERPH	AVA 16:10:1	veBPH)			•	. 47.4	86.7	1.68	n
				Bound	;	90 30	67	2.38	3.42	3.40	? ?	2.25	5.58	n
240	262053	H98812	3		7.69	07:07		2.89	3.39	3.14	X T		3.72	n
9	741067	AA478436	•nod	>	17.50	57.76	, ,	2.64	¥.,4	2.28	8.9	3 5	4.27	n
} ;	400178	AA121266	brain		35	11.48	· •	200	197	3.24	2.6 4	70.7	į	67
5 5	208741	HA3175		>	0.0	0.30	ກ (204	2.14	2.13	6.36	8 6		•7
707		20000	solo-		3.63	11.85	n	5 6	234	1.93	3.05	3.65	9 0	
33	110014	Weetee	5		0.50	1.62	ဗ	2.00	4	3.60	1.43	76.9	90.0	, •
ž	416092	Weses	•		0.55	1.79	е	1.18	<u> </u>	9 5	2.95	5.14	7.79	n :
255	22138	H92525	•		800	0.17	ო	3.24	9 C	\$ ¥	1.74	3.08	1.09	r)
5 2	223274	H86518	•		•	3.78	60	6.67	2.69	2 9	980	2.05	5.3¢	n
257	44387	H06525	brain			80	က	2.49	3.52	81.2 	3	3.80	2.03	60
258	110787	T90639	prain		67.0	86.5	. 11	1.59	4.12	2.42		3.17	2.10	Ð
259	121948	197762	prein) ;	3 9	· e:	4.87	5.06	12	2 3	2 00	4.45	60
280	782812	AA48251	brain		8.1	2 9	, er	1.90	4.01	2.67	97.0	27.6	3.25	60
28.	145513	R77718	s kin		0.67	2.10		2.30	5.27	1,85	3.63	i	58.0	0
•	435319		Pone		6.1	19.70	, c	5.5	2.24	3.47	9.80	07.0	2	n
707	90000		rojos	Maybe	.	4.52	n	9 8	921	3.78	3.20	2.11	1 000	
253	20000			•	0.36	1.16	က	2.00	2	2.43	2.00	3.26	18.2	, •
ž	18/08/			March	3.18	10.13	က	89.4	3 3	2.60	5.88	1 .08	80 60 60 60 60 60 60 60 60 60 60 60 60 60	,
202	416374		Done	anayan Aire	202	6.43	က	2.60	19.5	9	80.8	3.23	0.45	י מ
9 9	1422794	•		(90.0	က	1,47	4.27	<u> </u>	998	3.77	2.82	n
267	137017		•uoq		3 6	990	က	2.32	3.85	S. 5	9	1.72	3.	e
268	267634		breast		7 5	47 79	60	4.18	4.57	2.43	<u> </u>	2.73	4.79	n
269	433253		breast			32.56	ю	3.45	3.26	2.34	* C *	3.88	2.19	က
270	1475734		Neuron	∢ 2	8	4.60	. n	1.48	6.62	3.36	8 5	2.11	2.80	n
777	268876	N26026	•		- 1 - 1	7 49	· 67	3.66	4.92	99. 10.	3	3.5	4.13	77
222	79032	T61899	poue	>	7.61		· en	2.45	3.57	2.85	8	88.4	3.37	n
273	771023	AA427978	•	>	1,12		, es	1.08	2.45	1.78	t i	5.53	2,34	n
274	811101		brain		2.17	9.75) e	3.6	3.31	2.15	121	\$ F	127	n
278	270127	NZ7835	•		9 1	5.7	· «	3,50	2.77	3 .6	9.73	S.F.	2.39	n
276	448117	AA702480		¥	2.65	0.10		2.06	2.12	3.31	90.0	1.	4.45	
77.	796674		165		0.48		. "	1.65	4.32	2:32	3.80	7	3.82	n
278	431296						. 67	2.21	3.82		3 3	2.28	3.78	n
279	214624		prostate		7.82	20.0	. 67	4.39	4.19	1.80		240	1.91	n
280	743023	AA400596				3. °	, en	1.01	1.73	3.23	6.6	3	3.69	n
281	80050			₹	12.72	86.95	, en	2.12	4.97	1.27	,	Ę	3.08	n
282	1500162	2 AA886739	kidney	₹ Ž			, es	2.45	3.22	5.41	7.75	2.70	<u>;</u>	n
283	268234	N29992	Buni		2 3		, en	2.89	4.10	3.83	3.02	3.65	1.08	n
787	199623	R96561	•		6.9	1 2 2	, ea	2.41	3.80	2:88	3 7	7	1.32	ಣ
285	41305	R56677		Š	3 5	3	. 60	3.06	5.11	2.63		35.5	2.42	n
286	795555	5 AA459679	colon		3 !	,	, es	1.77	4 .00	2.37	70.4	88 7	1.97	n
287	416808	0 W86779	•		6.0		. 65	1.70	2.11	3.21	6.10	7 83	2.18	n
288	242698	H94262	•		5.7 5.1	800	. "	2.52	0.38	4.22	3.0	800	5,43	n
289	364352	2 AA022496	poue		3 3	9 6	· e7	1.60	3.79	1.68	77.7			
8	244050	N34042	•		80.0	5	,							
i								7						

Table 2-2: T3NO vs. BPH: Gene upregulated 5 fold or higher in at least 2 tumor samples or 3 fold or higher in at least 3 tumor Samples.

Order	IMAGE	Gen Bank Accession	Tissue	Secreted/Me mbrane	AveBPH	AveTumor	Fold(AveTumor/A veBPH)	Fold(MPM120/A	Fold(AveTumor/A Fold(MPM120/A Fold(MPM121/A Fold(MPM122/A Fold(MPM122/A Fold(MPM122/A Fold(MPM122/A # of Tumors veBPH) veBPH) veBPH) veBPH) Samples	Fold(MPM124/A veBPH)	Fold(MPM122/A veBPh)	Fold(MPM123/A F	Fold(MPM125/A 1 VeBPH)	of Tumors Samples
		Number		Bound		;	•	ç	1	330	3.80	2.68	2.56	n
291	544639	AA074677	skin		Ę.	3.15	9	200		284	97	2.83	0.82	n
282	852520	AA663058			7.88	\$3.5¢	r)	85.5	5			1.07	11.7	n
282	797000	AA463500	toell		0.11	0.33	က	2	3.21	8.		63	2.20	•
ă	1403252	AA886870	orostate	¥	18.18	54.08	eo	4.82	3.25	2.81	97.5	4 4		. •
ğ	477416	AA702335	brain	¥.	1.15	3.41	က	5.58	4.1	2.12	5	2 (.	
3	0/63477	2002200	i .	W.	15.	4.56	ю	2.92	3.27	£.53	8:	8	2 1	3 (
	9797	/88//97	•	2		0.65	n	1.35	3.30	3.86	2.74	3.80	2.47	r)
787	10.00	SDICH.	•		* :) F	; er	5.57	3.38	1.71	2.34	0.80	3.84	n
8	1400837	AA863449			2 5	3 5	• •	2.76	3.98	2.31	3.06	2.04	3.43	n
8	20183	H48278	•		3 5	* .	. "	3.82	7.88	3.41	1.10	1.08	0.29	n
8	263769	H00694	•		6/:0	2. 7	, ,	800	3.14	64.0	0.21	4.4	277	n
8	427930	AA001976			900	†	, ,	0000	4.92	3.17	2.07	3.86	1.25	••
305	307244	N95187	hear		8	5 !	, i		3.18	1.43	200	4,41	<u>‡</u>	n
g	810843	AA458959	brain		9 9	0.17	m (40°	? ;	3.20	1.57	3.43	4.63	n
ğ	1460820	AAB89785	pone	¥	3.62	10.49	m	2,35			40	2.31	2.77	•
88	26259	R20547	prostate	>	0.87	2.50	က	3.08	3.41	90.0	2 5	8	6.23	60
8	386154	AA062813	poue		0.0	0.03	m	3.87	3.77	8.	3 5	3 6	į	
202	282104	N51498	brain		0.83	2.66	es es	2.66	4.75	3.79	Ze: 1	2 .	7.	
Ş	344308	W70147	,	>	0,46	1.31	ຄ	3.18	3,47	5.50	1.78	5	• •	· •
} }	45525	108724	eteterre	· >	2.19	6.17	ю	2.20	3.24	3.18	3.52	2:33	2.7	, .
3 3		24,000		•	8	0.11	n	2.56	4.53	3.0	2,48	8. 8.	8	? (
2 :	267406	Naches Naches		>	9	4.40	ຄ	3.68	3.24	. 2.59	3.41	1.52	2.19	n (
5	201101	707C7N	9	-	8	1.78	- г	3.73	4.67	3.45	1.24	8.	1.53	י מ
212	001/0/	3			} ;		· e:	2.36	4.23	3.40	3.07	2.83	0.52	n
313	204122	H61886	•		3 5	, ,	, e	5.37	0.00	1.38	1.18	3.6	3.82	n
314	638545	AA489791			7	* ::	· •	7	95.0	3.30	2.89	4.45	4.47	n
315	809694	AA454702	brain		20:0	8 6	? (3.70	2.31	1.13	3.56	3.32	ဗ
316	126234	R06362	•		8	1970	? (2 7	9.40	191	3.26	3.22	2.72	n
317	280407	N51563	nuspecified	¥	0.97	2.62	.	10.1) r	÷	3,46	3.11	4.17	n
318	758284	AA404231	placents		0.78	2.08	m	797	10.4	2 .	75.	1.07	5:00	n
319	150314	H00617	tcell		2.18	5.81	ო	3.69	Q. !	67.0		8	3.97	e
320	288936	N62696	٠		0:0 *	0.11	ဂ	1.59	3.47	9, 50	3 5		805	. es
32	757440	AA437226			90.0	0.13	60	0.21	4.10	1.76	70'4		9	. स
322	176554	H45289	•	¥N.	0.87	2.28	ဗ	1.89	2.02	3.72	3.33	00.0	3 5	. e.
323	296057	08009X	hear		99:0	1.79	က	2.03	3.01	7.2	S :		3	
324	755299	AA496359			1.69	4.39	6	4.47	0.52	8	93.6 6	2.5	27.	• •
325	725321	AA201740	breast		0.87	2.27	೮	3,21	3.07	3,61	2:30	70'1	2	
328	1467831	AA883431		ž	1.17	3.02	6	1.16	1.22	4.32	0.72	33.55 1.55 1.55 1.55 1.55 1.55 1.55 1.55	70.4	· •
2	120007	TORIGI	Meridia		0.00	45,	ю	2.09	2.01	* .	3.86	3.01	180	,
, K	159455	H15746	prain		0.02	9.0	ь	0.62	3.70	2.31	3 3	3,72	F. 6.) (
2	AAAAAA	4 4 9 4 9 5 0 0		***	6.57	16.82	m	1.39	3.08	ğ	2.32	3.41	77.	•
	76776	4.40729	, }	<u> </u>	000	90'0	n	3.92	2.20	3.37	0.77	0.24	28 ,	n
3 :	450613	77451 144V		4/2	0.74	1.89	e	1.17	98.0	0.92	3.00	4.79	‡	•
ā	1000	7C4790VV	• }	Š	3 5	2	• •	4.17	‡ ‡	99.0	1.17	4.50	0:30	n
ñ	42343	W68141	Poop		70.0	Š	•							

Table 2-2: T3NO vs. BPH: Gene upregulated 5 rold or higher in at least 3 tumor Samples.

		IMAGE	Gen Bank		Secreted/Me mbrane	AveBPH	AveTumor	Fold(AveTumor/A Fold(MPM120/A Fold(MPM121/A Fold(MPM124/A Fold(MPM120/A	Fold(MPM120/A	Fold(MPM121/A	veBPH)	veBPh)	veBPH)	(MBPH)	Samples
1,00,00,00,00,00,00,00,00,00,00,00,00,00	=	Clone 1D		Distribution	Bound					0	3,58	1.37	1.19	1.85	က
		*******	997707	ri acra		5	2.65	ဗ	3.31	3.65	9 6	25	3.28	1.79	60
		200	AA131400	- ·	A design	17.73	11.11	ຕ	3.17	4.60	0.70	<u> </u>	0.31	3.14	63
255051 WAZZEN DOWN MAZZEN DOWN DOWN DOWN DOWN DOWN DOWN DOWN DOW		784130	AA4 32075	prostate	Mayod		800	က	3.68	3.87	2.82	2 6	1.37	3.87	60
244599 MYSTAN Model Office 1 0		325370	W52208	pou		3 6	40	es	3.07	3.96	0.85	3.		2	n
944589 NATESTY Imp 1557 14459 1459 1459 1450		288663	N62394	kidney		ا رو در	2 6	, ,	0.90	3.52	41.0	3.08	70.7	¥. +	•
CASEAL FRIENCE 115 279 2 35 319 112 116 379 2 35 319 112 118 119 22 225 319 112 118 119 319 319 118 319 <th< td=""><td></td><td>344359</td><td>W73377</td><td>- Bunt</td><td></td><td>0.07</td><td>5. 5.</td><td></td><td>28</td><td>3.57</td><td>3.34</td><td>3,40</td><td>2</td><td>3 :</td><td>•</td></th<>		344359	W73377	- Bunt		0.07	5. 5.		28	3.57	3.34	3,40	2	3 :	•
CASISTIAN ACCORDINATION CASISTIAN	_	63301	R16241	•		1.15	2.79	81	200	10	1.82	1.65	1.06	3.55	,
200212 Columnation Columnation <t< td=""><td></td><td>430824</td><td>A A 0.008.03</td><td>•</td><td></td><td>0.14</td><td>0.35</td><td>7</td><td>3.10</td><td>• G</td><td>28</td><td>4.12</td><td>0.85</td><td>2.32</td><td>n</td></t<>		430824	A A 0.008.03	•		0.14	0.35	7	3.10	• G	28	4.12	0.85	2.32	n
2005013 ACTION CONTINUE OLD OF CONTINUE		C70874	20000	,		0.62	1.48	2	3.61	37.	3 8	1.52	3.40	3.14	60
452080 AVATOR NATION NATION<	_	747907	CCCC	•		000	0.20	7	0.60	4.21	3 (3.01	1.39	eo
45000 AVANTIS Intent 2009 6684 2 249 310 0.048 337 0.028 3587 AVANTIS Intent 2009 6684 2 249 310 0.048 337 0.028 3587 AVANTIS Intent 2009 6049 0.049 2 4.12 3.40 0.04 3.09 0.048 337 0.028 3587 0.028 AVANTIS Intent 2009 0.049 0.009 2 4.12 3.40 0.04 1.00 0.04		255897	N27366			8	2.03	7	1,29	3.24	1.70	5	£ 0	3.18	63
586779 ALTASSAPO Decided as a composition of the co		452068	AA707125	Teet Teet		8 5	3 97		2.43	3.80	0.16	3.5/	3 3	405	n
1,0,0,0,0,0,0,0,0,0,0,0,0,0,0,0,0,0,0,0	_	585779	AA135870	prostate		20.69	40.04		1 20	3.87	0.24	3.30	\$ 70	8 6	•
SENORIS AMERISOR Death 0.26 0.76 1.7 1.8 0.74 4.24 SENORIS AMERISOR Death 0.06 0.06 0.06 0.76 1.6 0.74 1.76 0.74 1.76 0.74 1.76 0.76 1.76 0.77 1.77 0.77 1.72 0.77 1.19 0.44 0.77 0.48 0.44 0.48 0.44 0.48 0.44 0.48 0.44 0.48 0.44 0.48 0.44 0.48 0.44 0.48 0.44 0.48 0.44 0.48 0.44 0.48 0.44 0.44 0.44 0.44 0.44 0.44		128506	R10154	prostate		6	0.00	•		376	0.42	3.04	0.88	28.0	, (
CORRADIA ALTISTICAL STATES CORPORABILISM O.01 0.14		870078	AA487505	postu		0.38	0.76	74	3.16	2 6	2	0.74	42.48	32.88	N
CATACORDIAL ALL AND COLOR ALL ALL ALL AND COLOR ALL ALL ALL ALL ALL ALL ALL ALL ALL AL	_		2000	all affects		0.01	0.18	ŧ	0.74	t :		78.7	34.78	5.82	~
1777047 H1680265 NAME of the control of		2000	791961AA			8	0.50	60	1.80	0.16	90'7	93.0	11.1	2.44	~
137894 RECOLOR Suith Color C		50787	H16833			2	2.25	80	8.43	33.82	0.3/	3 :	** 62	1.71	~
666563 AAA608222 breast 119 0.74 0.75 0.75 0.25 1.05 0.40		137984	R63085	s kin		•		œ	1,46	0.70	1.10	?	1 :		7
		856585	AA869222	bresst		0.12	S	,	7.32	29.35	1.05	0.49	9	3. ·	•
61323 TOOTIT Y 0.47 3.19 7 1.55 2.288 0.657 1.63 1.01 820202 H.23222 1.68 7 1.67 0.12 0.12 0.02 31.39 42006 H.2022 1.69 0.22 1.59 0.71 0.12 0.02 1.07 0.12 0.13 0.13 0.13 0.13 0.13 0.13 0.13 0.13 0.13 0.13 0.13 0.13 0.14 0.13 0.14		151477	H02837	skin		4.10	8.16	- 1	£ 33	19.31	2.50	<u>7</u>	0. 68	3 :	
SSCORE HZZEZZZZZZZZZZZZZZZZZZZZZZZZZZZZZZZZZZ		81331	T60111		>	0.47	3.10	,	3 5	22 88	0.57	3.1	1 .9	0.38	•
BORDATION PARTICION CONTINUE NATIONAL CONTI		#202#	L22032	•		0.25	1.86	4	13.07	3 6	0.12	0.52	31.38	5.36	~
4000000 AMADOS (MARCO) AMADOS (MARCO) AMADOS (MARCO) AMADOS (MARCO) AMADOS (MARCO) MARCO)		2000	100000	poloc		90.0	0.53	ø	1.07	7.7		1.35	18.46	16.17	~
24440 NALOSTION 0.006 0.32 5 6.88 2.35 1.77 2.77 11.30 754368 AALOSDIAZ bone 8.26 44.86 5 0.04 1.57 1.77 2.77 11.30 754368 AALOSDIAZ bone 8.26 44.86 5 0.04 1.67 0.02 2.06 757168 AALOSDIAZ bone 1.00 0.04 5 7.84 1.47 2.77 1.47 0.04 0.05 0.06 0.05 0.06 0.05 0.06 0.05 0.06 0.05 0.06 0.05 0.06 0.05 0.06 0.05 0.06 0.05 0.06		70.00	AA000403	5		0.22	1.39	80	0.88	14.0	8 8	0.55	6.1	20.60	~
244368 AAA3562 Done 6.25 4.496 5 2.04 2.77 1.17	_	975	Od I CONTA	Ď		90.0	0.32	¥O	8.98	2.33	<u>.</u>	74.6	11.30	12.03	7
754536 AAA40512 Done 0.7 1,51 0.84 A.A25145 0.64 1,51 0.84 A.A25145 0.64 1,51 0.84 0.61 0.7 1,61 0.84 0.61 0.64 0.61 0.43 0.62 0.62 0.61 0.64		264448	NZTZSS	• ,		8.25	44.96	NO.	2.04	2.77	37		2.06	5.30	~
7871484 AAA45950 Namon 1825 80.08 5 7.86 14.43 2.22 1.41 0.0 47355 AA443950 Namon 18.26 80.08 5 14.30 0.10 0.55 0.22 9.86 47355 AA196125 besits N/A 3.49 16.28 4 5.72 2.44 11.80 1.20 2.30 377895 AA77706 lesits N/A 0.46 1.95 4 1.17 7.49 1.07 1.48 0.06 79703 T63961 lung 0.04 0.71 4 1.17 7.49 1.47 1.47 1.49 0.00	_	754358	AA436142	euog :		5	0.16	NO.	0.47	1,51	3 .	4 ,	190	2.40	8
665235 AA136125 Deesat 19.20 0.10 0.46 5 14.30 0.10 0.65 0.22 0.22 0.23 0.22 0.23 0.23 0.23 0.23 0.23 0.23 0.23 0.23 0.23 0.23 0.23 0.23 0.23 0.23 0.24 1.14 1.45 1.23 2.30 0.23 0.24 1.45 1.23 2.30 0.23 0.23 0.24 1.45 1.45 1.44 1.47 1.47 1.44 1.45 1.44 1.45 0.05		757165	AA443950	Neuron		35.85	80 08	ĸ	7.86	14.43	2.52	*	9	2.80	~
47355 H10881 NA 349 15.28 4 5.72 2.44 11.69 1.29 4.20 377888 AA77708 leafs NA 3.46 16.28 4 6.41 14.50 1.23 1.41 1.47 745072 AA26275 NA 0.16 0.75 4 1.17 7.46 1.07 1.46 0.06 745072 Te3961 Color 0.16 0.72 4 1.17 7.46 1.07 1.47 0.06 1380016 AA855158 Color 0.20 4 1.17 7.46 1.07 1.46 0.02 284331 N71861 - 0.02 0.07 4 1.12 0.65 2.67 1.86 103120 AA60982 - 0.02 0.07 4 1.17 0.36 0.57 0.57 0.57 0.57 0.56 0.56 0.50 0.57 0.56 0.56 0.56 0.57 0.56 0.56		565235	AA136125	Dreast			870	· va	14.30	0.10	0.55	77	9 6	2.82	~
377898 AA777098 leatis N/A 3.48 1920 4 641 14.50 1.27 1.41 1.45 1.47 1.47 1.48 1.47 1.47 1.48 1.47 1.48 1.49 1.47 1.48	_	47355	H10981			2 9	3 5	•	5.72	2,44	11.68	1.28	χ.,	180	7
745072 AA626275 AA626275 NAA 0.45 1.73 7.49 1.07 1.48 0.00 79763 75963 75964 1.09 0.17 4 11.10 10.86 0.50 0.52 0.21 138018 AA655158 colon 0.05 0.20 4 11.10 10.86 0.50 0.57 1.88 28333 N5141 . 0.18 0.72 4 1.22 10.53 5.68 2.67 1.88 284831 N71861 . 0.02 0.07 4 1.17 0.39 5.73 1.64 0.35 1046992 AA60992 . 0.03 0.11 4 1.07 0.39 5.96 2.96 2.67 0.36 1046992 AA778536 endothelial NA 0.14 0.57 4 0.51 2.96 2.96 6.06 404699 AA6189 noisi 1.18 4 1.87 2.96 2.96 <	_	377898	AA777098	testis	Ş		97.01		6.41	14.50	1.23	¥.	/ v :1	2 7	. ~
79763 T63961 lung 0.16 0.71 4 11.10 10.86 0.50 0.52 0.21 1389018 AA865158 colon 0.05 0.20 4 11.2 10.63 5.86 2.67 1.86 283335 N51441 0.18 0.72 4 1.22 10.63 0.67 0.82 284831 N71861 0.02 0.07 4 1.17 0.36 5.13 1.64 0.36 1031203 AA60962 0.03 0.11 4 1.17 0.36 5.13 1.64 0.36 103120 AA682274 breast N/A 10.27 38.51 4 1.87 2.96 2.96 2.21 1.03 461699 AA682274 breast N/A 10.27 38.51 4 1.87 2.97 2.86 2.21 1.03 461699 AA682274 breast N/A 1.027 38.51 4		745072	AA626275		₹ Ž	o 5	SK:	• •	1.17	7.49	1.07	 84.	0:08	ţ ;	
1389018 AA855158 colon 0.05 0.20 4 1.22 1.053 5.68 2.67 1.88 283335 N51441 . 0.18 0.72 4 1.22 10.53 5.68 2.67 0.87 0.82 284831 N71861 . 0.02 0.07 4 1.77 0.36 5.13 1.64 0.36 1031003 AA60082 . 0.03 0.11 4 1.073 5.96 2.96 2.22 1.03 1041990 AA682274 breast N/A 10.27 38.51 4 1.87 2.96 2.96 2.96 6.06 461890 AAA018980 brain 0.31 1.18 4 1.87 2.97 2.86 2.21 1.03 56022 AA118980 brain 1.07 4.00 4 2.76 5.27 1.60 11.37 0.21 12216 AA118980 brain 1.07 4.00 4 <		79763	T63961	Bung		0.1 6	0.71	•		10.86	0.50	0.52	0.21	9.1	• •
283335 N51441 . 0.18 0.72 4 1.24 8.36 0.83 0.57 0.92 284831 N71861 . 0.02 0.07 4 2.88 8.36 0.13 1,64 0.36 1C031203 AA606082 . 0.03 0.11 4 1,17 0.36 5.96 2.96 2.22 1.03 1C045060 AA778636 Andothelial N/A 0.14 0.67 4 0.81 2.86 2.96 2.96 2.22 1.03 461690 AA778636 Andothelial N/A 10.27 38.51 4 0.81 2.86 2.96 2.96 2.22 1.03 461690 AA682274 breast N/A 10.27 38.51 4 1.87 2.86 2.96 2.26 6.06 552226 AA018980 brain 1.07 4.00 4 2.76 5.27 1.54 0.20 248806 N53214 b	_	1389018	AA855158	colon		0.05	0.20	•	2 .	10.53	5.68	2.67	1.88	3.	•
284831 N71861 0.02 0.07 4 2.08 6.13 1.64 0.36 1C01203 AA609682 0.03 0.11 4 1.17 0.38 5.13 1.64 0.36 1C04896 AA778636 andothelial N/A 0.14 0.67 4 0.81 2.86 2.96 2.22 1.03 461696 AA778636 andothelial N/A 10.27 38.51 4 0.81 2.86 2.96 2.22 1.03 461696 AA682274 breast N/A 10.27 38.51 4 0.81 2.87 2.86 2.21 6.06 56022 H17826 adramal 1.07 4.00 4 2.76 5.27 1.50 0.71 122176 T98616 b 0.03 0.11 4 6.45 1.22 1.34 0.20 248808 N53214 b 0.61 2.24 4 2.03 2.16		283335	N51441	٠		0.18	0.72	₹	4 6	3.0	0.83	0.57	0.82	10.57	7
1001200 Aveoges		284031	N71861	•		0.02	0.07	₹	2.08	3 6	5.13	3.	0.36	15.08	7
1001AU Accordance Accordance<		,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,		•		0.03	0.1	→	1.17	S		226	8.	99:0	7
461696 AA682274 breast N/A 10.27 38.51 4 0.81 2.58 2.78 2.21 5.65 50602 H17626 adrenal 0.31 1.18 4 1.87 2.76 5.27 1.60 11.37 0.71 362026 AA016980 brain 1.07 4.00 4 2.76 5.27 1.60 11.37 0.71 122176 T98615 . 0.03 0.11 4 0.65 8.21 0.34 1.94 0.20 246808 N53214 b 0.56 2.06 4 6.45 1.20 1.20 0.84 0.52 191866 H40350 . 0.61 2.24 4 7.00 9.56 2.16 0.82 0.72 155072 R71393 . 0.30 1.07 4 7.00 9.56 2.16 0.72		30000	•	alledoboe	V	0.1	0.57	•	10.73	8	B 6	845	909	2 ,	~
451699 AA682274 breast NA 1.18 4 1.67 2.97 2.88 2.21 0.71 5.692 H17626 admenal 0.31 1.18 4 1.67 5.27 1.60 11.37 0.71 0.71 5.692 AA018990 brain 1.07 4.00 4 2.78 8.21 0.34 1.94 0.20 1.22178 T99616 . 0.65 2.06 4 6.45 12.29 1.20 0.84 0.52 2.46608 N53214 b 0.65 2.06 4 2.33 6.76 2.63 2.89 5.39 15672 R71393 . 0.30 1.07 4 7.00 9.56 2.16 0.92 0.72	_	2000	•			40.27	38.51	•	0.81	2.56	94:7		X A X	6.87	~
50892 H17626 adrenal 0.51 1.37 0.71 362926 AA018890 brain 1.07 4.00 4 2.76 5.27 1.94 0.20 122176 T98616 - 0.03 0.11 4 0.65 8.21 0.34 1.94 0.20 246808 N53214 b 0.66 2.06 4 6.45 12.29 1.20 0.84 0.52 191866 H40350 - 0.61 2.24 4 7.00 9.56 2.16 0.92 0.72 155072 R71393 - 0.30 1.07 4 7.00 9.56 2.16 0.92 0.72		461699	AA682274	Dreast.	C	2 2	1 18	•	1.87	2.97	2.88	27	3 6	17.0	~
362026 AA018890 brain 1.07 7.00 9.56 8.21 0.34 1.94 0.20 1.20 1.20 1.20 0.84 0.52 246808 N53214 b 0.56 2.06 4 2.33 6.76 2.63 2.98 6.39 1500 0.81 0.32 0.72 1500 0.81 0.82 0.72	•	2000	H17626	equene e		9 6	. 4	4	2.78	5.27	9.	11.3	- 6	10.81	8
122176 T98616		362926	AA018980	brain		<u> </u>		•	0.65	8.21	0.34	ă.	3 6	A B C	7
246806 N53214 b 0.56 2.00 4 2.33 6.76 2.63 2.98 6.39 191866 H40350 - 0.61 2.24 4 7.00 9.56 2.16 0.82 0.72 155072 R71393 - 0.30 1.07 4 7.00	_	122178	T98615	•		3 1	. 8		6.45	12.29	1.20	9.0 7	70.	3 8	•
191866 H40350 - 0.61 2.24 - 7.00 9.56 2.16 0.92 0.72 155072 R71393 - 0.30 1.07 4 7.00	~	246808	N53214	۵		9 0 0	8 7	• •	2.33	6.76	2.63	2.98	5.38		
156072 R71363 . 0.30 1.07 4	•	191866	H40350	٠		9.0	2.24	• -	200	9.56	2.16	0.92	0.72	3	•
		155072	R71393	•		9 9 9	1.07	•	<u>.</u>						

ž
8 ample 8 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2
Fold(MPM126// veBPH) 6.75 1.08 2.39 1.86 0.69 5.09 2.25 1.06 1.06 1.28 0.70 8.46 5.38
Folid(MPM120/A) Folid(MPM121/A) Folid(MPM122/A) Folid(MPM1
Fold(MPM122/A weBPh) 7.62 1.03 1.98 7.16 6.92 2.76 8.52 2.76 5.55 1.53 1.33 1.33 5.44 5.17 5.44 5.45
Poid(MPM124/A veBPH) 2.17 1.08 7.88 2.71 1.89 1.89 1.44 2.75 1.59 5.26 5.26 5.26 5.26 5.26 5.26 5.26 5.26
Fold(MPM121/A veBPH) 2.12 1.08 0.37 1.16 5.29 6.18 6.18 7.10 7.10 7.10 7.10 7.10 7.10 7.10 7.10 7.10 7.10 7.10 7.10 7.10 7.10 7.10 7.10
Pold(MPM120/A VeBPH) 1.28 6.85 6.17 1.42 2.20 6.85 1.38 0.69 5.71 5.36 2.60 2.59 0.69 0.69 0.73
Fold(AveTumor/A veBPH) 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4
3.94 0.03 0.10 9.83 1.82 0.21 29.44 0.71 4.54 5.67 3.90 0.10 0.32
Ave BPH 1.10 0.01 0.03 2.85 0.05 0.07 9.12 1.45 1.45 1.27 2.07 0.04 0.04
Secreted/Me mbrane Bound N/A N/A
Tissue Distribution bone breast muscle breast lung lung lung testis brain prostate lung
Gen Bank Accession Number AA134862 N69675 AA63239 AA63239 AA6315 H15366 AA023028 AA62801 AA623034 H16743 W83776 AA453518
IMAGE Clone ID 502436 283689 510136 878284 289025 94713 796646 49315 1684274 36455 743688 60114 342164 785407

order.

Table 2-3: T3NO vs. BPH: Genes upregulated 10 told of higher in at least 3 tumor samples or 5 fold or higher in at least 3 tumor samples.

order.		Gen Bank		Gen Bank Secreted/Me			Fold(AveTumor/Av Fold(MPM120/A	Fold(MPM120/A	Fold(MPM121/A Fold(MPM124/A	Fold(MPM124/A	Fold(MPM1220 veBPh)	VeBPh) VeBPH) VeBPH)	veBPH)	Samples
	MAGE	Accession	Dietribution	mbrane	AveBPH	Avelumor	•8PH)	(HARPA)			63 43	22.54	10.36	•
	500	Number		Bound	8	133.47	67.99	149.63	165.70	87 ! 	10.30 10.00	53.59	35.90	•
	788180	AA453310	prostate		3 6	94	47.98	65.24	24.62	19:11	80.8	41.24	19.50	•
	648037	AA198979	prostate		8.30	3	30.37	31.27	50.07	34.87	27.0	45.42	20.45	•
	284701	N64840	prostate	>	2.41	2.5	30.08	38.52	11.57	48.94	10.01	21.41	63.61	•
	450049	AA703396	prostate	Š	2.53	\$;	23 40	11.34	8.80	10.00	77.61		13.29	•
	245768	N55266	•		0.01	0.21	\$ F.7.7	22.18	20.64	22.40	16.57	ē ;	1.3	•
	2003	N50880	7		24.06	423.65	70.11	27.00	41.52	9.28	10.57	C. 6		•
.	20107	036061	į	>	5. 2.	44.38	17.34	97.67	78.57	6.15	9.24	10.23	QX.G	•
~	130000	ancer!	4	· >	1.48	25.48	17.23	30.48	3 7	11.00	10.01	47.00	90.0	•
	133130	R28397		-	0.0	0.12	11.81	9.28	\$	82.0	7.12	8.55	19.24	6
•	179753	H51549	9 00		202	19.56	9.43	6.78	6.32	8 8	25.	12.71	6.41	•
9	51433	H20747	•		9 6	8	7.67	5.89	6.60	5.6	3 6	19.49	9 .01	w
=	195975	R91398	•		9.0	8 8	12.94	15.20	2,45	±. ₹.	8 2	13.28	1.75	*0
12	287528	N62096			9	8 5	10.47	7.91	24.39	10.24	5.2/	4.27	15.03	10
£.	259374	N31952	prostate		1.29	13.32	4,04	7.99	15.61	11.93	6 .14		7.08	80
: 3	1030855	AA621761		>	7	12.57	10.15 4 85	5.69	10.96	6.72	12.64	3 5	8.20	¥O
. 1	1031599	AA609485	•		0.27	2.07	9.0	3.24	10.05	6.80	7.18	0.0	14.57	10
	21931	172543	breast		0.29	/8.T	0.0 3.4	6.28	4.94	5.38	5.13	570	80.6	10
-	825859	AA504784	٠	K/N	<u>6.</u>	VE.21	A 82	7,23	6.80	5.30	86.28		19.00	*
	130136	R20813	٠		\$	Q.0	34.18	18.89	1.69	59.34	105.24	8.5	1.26	*
•	1468260	AA884926	•	₹Z	0.65	22.36	2 5	29.78	48.11	29.58	0.86	79:07	27	*
. 8	743828	AA634379	pone	ΚX	0.58 83	12.90	78.97	9.65	9.81	<u>.</u> Ž	57.03	3÷.61	40.5	*
7	41647	R52794	brain		0.40	1	5.00	9.62	26.16	14.54	17.70	77.	72.5	*
3	784168	AA432103	prostate		15.97	194.72	12.18	10.87	18.00	7.67	13.10	7 :		•
1 5	193923	R83407	brain	>	17.59	158.73	70.0	13.49	10.45	13.83	13.97	8.5	3 2	4
;	487165	AA045074			0.0	000	9 F	13.54	16.10	7.85	4.72	g !		•
; ;	306066	N91003		>	0.80	3 .	0.70	70,0	10.11	7.14	10.04	4.07	- 6	. 4
3 :	202217	08197N	pooq		90.0	0.42	6.73	1 69 1 69	10.38	6.62	10.46	7	3 7	• ◄
e !	07277	00000	prostate		0.19	1.27	6.55	8 8	10.82	5.48	10.02	<u>4</u>	3.81	•
3 8	887687	AAROOROS		¥X	0.38	2.38	46.0	9 9	80	10.11	2.19	1.74	10.42	• •
2	4068473	*****	Ē		11.61	70.00	6.03	9.00	44.77	5.12	5.02	1.51	2.6/	•
20	7/10001				0.95	5.55	5.84	8.83		6.21	7.48	3.67	8.43 5.43	•
8	182383	De12-00			0.02	0.14	5.84	3.19	10.0	2	6.43	5.07	2.77	•
5	195034	R85764		4/2	, ,	1.88	5.48	2.60	0.47		1.30	0.26	5.98	₹
g	461541	AA705258		<u>{</u> }	1.76	8.87	5.04	6.20	7.0	3.6	1.52	20.0	9.17	*
ន	787938	AA452278	g.	-	0.02	0.10	5.01	5.74	6.67	S. C.	85	5.02	1.7.1	4
ま	754034	AA479058			240	2.07	4.38	5.34	6.4 6	8.00	į	3.32	7.01	6
3	195387				7	46.10	13.39	11.55	50.28	3.85	20.6	06.0	1.22	n
8	83868	•				8	12.85	43.16	25.22	F0.6		18.31	0.82	•
3	281039		•		2 2	0.46	12.37	46.34	6.67	1.83	7 5	48.21	5.35	•
8	490718	•			\$ 8	27.62	11.81	2.78	4,13	1.86	70.0	•		
8	744944	AA625890	or or other		3	- F								

Table 2-3: T3NO vs. BPH: Genes upregulated 10 fold or higher in at least 3 tumor samples.

Order		Gen Bank	i	Secreted/Me			Fold(AveTumor/Av	Fold(MPM120/A	Fold(MPM121/A	cold(MPM124A	veBPh)	VeBPH)	Fold(AveTumor/Av Fold(MPM120/A Fold(MPM121/A Fold(MPM124/A Fold(MPM124/A Fold(MPM120/A	Samples
	IMAGE	Accession	Distribution	mbrane	AveBPH	Averumor	•BPH)	(HARA)			¥,	0.86	1.67	n
	Clore	Number		Bound	i	;	10.89	20.89	31.93	7.51	67		19.1	•
\$	287728	N59158	•	>	0.51		8 6	22.49	23.74	3.29	9.26	707		•
: =	1034473	AA779728	prostate	¥X	0.13	1.36	10.50	10 16	21.53	1.58	8.31	4	2 4	, et
;	7678376	AABA7007	breast		0.08	0.69	0.50 0.50	10.04	0.73	2.20	24.39	4.57	2.6	•
; ;	200456	44043260			0.0	0.11	8.02	10:01	A 48	4.03	15.83	2.66	4.48	, ,
3	200	2000			1.71	11.25	6.59	as:	? e	72.63	2.12	4.38	6.18	,
\$	96 96 96 96	172336	Brod		78.0	3.71	6.48	8.59	3.80	976	980	2.35	3.29	n
\$	268211	N29986	prostate	;		80.24	6.21	9.37	12.99	04.7	7	4.07	3.73	n
\$	593185	AA165678		>-		Ş	6.03	6.54	8.44	9. 1		2.69	8.45	60
!	428816	AA004681	٠		2 5	3 6	5.92	9.12	4.48	4.71	\$ 1		7.68	n
\$	811582	AA454597	poue	>	200 P	77.0	, A	1.13	4.65	4.73	r/.e	75	3.73	n
3	428436	AA004412	٠		0.53	31.5	 	5.69	4.98	0.83	11.93	*	2.12	•
8	795729	AA460291	pegipedeun		0.02	רנים י	9 9	88	14.24	8.62	0.72	8 3	27	n
2	712950	AA282273	1.09		5.83	32.98	6 1	3.24	6.98	2.95	6.31	10.8	2 5	• •7
; {	825785	AA505117	OVBIY		. .5	8.57	C:C	900	0.51	7.72	0.13	17.97		
3		200000	900		8	0.43	5.45	07.0		7.31	8.02	273	2.67	n (
3	204/60	AA146/30			0.75	3.90	5.23	3.83	7.0	2 6	908	2.79	3.31	n
3	491706	AA115259	Dream		5	5.43	5.10	3.85	6.61	78.0	8	7.48	8.28	n
8	241821	H93381		Ç Ž	<u> </u>	09.8	5.01	2.52	2.75	20. F	3 6	1.13	5.84	ю
8	782537	AA48484	tostis		7 :	200	78.7	7.68	12.12	2.17	3 :	878	5.11	0
25	277596	N57002	pone	>	2 :	ţ	7	2.53	5.09	3.89	9	e s	2.93	n
8	124128	R01348	•	>	1,25	0.0	8 4	5.02	7.58	4.08	5.94 4	3.5	202	n
2	134719	R28287	Color		0.73 0	9	2	5.23	10.10	4.41	5.26	91.T		•
3 8	814306	AA459318	prostate		9.65	45.42	L' -	970	4.83	6.38	7.51	. .3	90.7	. "
3 3	784273	A4446316			0.02	0.10	07.4	9 9	50.0	6.74	1.93	1.66	27.82	, «
5	2007	240070	,		0. 4.	0.67	4.62	69.G		523	1.01	1,63	89.4	, ·
8	3	0/0817	,		12.85	59.29	4.61	5.52	8 6		8.18	1.05	7.68	n
ន	260170	N32072	brosense		0.02	0.09	4.56	1.67	6.27	3.4	83	3.29	3.63	n
Z	277327	N57483			5	15.26	4.54	9.25	5.31		340	0.26	2.38	n
3	307249	N93438	tosus.		7.38	33.36	4.52	7.28	8.45	D (0	2	2.43	0.38	n
8	770086	AA430545	ciero		3 5	0.12	4.49	1.90	8.33	87.7	3 .	986	5.65	n
67	51620	H19446	•		3 3		4.38	2.17	5.43	7.7	8 .	72.	8.83	n
3	428267	AA004626	pegpedeun		- ·	2 6	75.7	0.98	7.98	1.80	57.6		6.79	n
8	278504	N66139	nerq		3 3	2 18	4.28	7.69	2.40	5.57	97.1	\$	6.16	n
2	205628	H57585	prostate		5 6	0.0	4.23	6.18	3.28	3.22	3 8	800	9.45	n
۲	265005	N21081	s kin		3 6	90	4.16	5.29	7.89	0.68	3 3	9	1.52	n
22	67397	T49325	testis		5 5	800	4.03	2.09	6.18	6.46	57.7	97.0	\$3.3	n
r	284624	N64796			8 6	24.0	786	0.28	7.63	0.28	G 1		8.10	n
7	882497	AA878604	pou		5 6		3.89	6.41	6.16	8 6.0	0.30 0.30	3	6.57	n
75	856174	AA630620			3 3		3.78	1.62	0.09	1.78	6.83	r ;		n
76	503824	AA133959	brain		0.12		7	5.28	5.92	7.33	. 69:	0.76	. e	•
-	208210	H65286	placents		0.80	4		88	1.13	5.78	5.41	51.1	60.0	
	325380	AA284285	spieen		0.0	0.03	0.40	7.0	1.13	1,13	14.34	13.4	2 3	
: ;	270017	N32514	•		0.0	0.23	29.82	£0'6	57 87	3.22	1.15	2.77	3.51	• •
2 1	1800KB	4 4 8 4 3 5 0 3	etataco		2 .	33.00	18.02	49.03	£	0.11	0.61	3.63	22.02	N
2 3	Page 1	300000V		>	070	3.10	15.42	64.92	77.1	;				
5	3338	2/15#4		r				8						

84 TO 10 10 10 10 10 10 10 10 10 10 10 10 10
Fold MPM126A VeBPH) 2.05 32.88 2.28 4.21 1.40 0.36 15.67 12.03 1.46
Feld(NPM123A veBPH) 44.95 42.46 1.07 0.58 0.58 1.01 1.01 1.47 11.30 0.21
Fold(MPM122/A veBPh) 28.47 0.74 1.88 1.28 1.64 1.63 2.85 1.35 2.85 2.85 2.85 2.85 2.85 0.52
Fold(MPM124/A veBPH) 4.77 1.86 4.51 1.81 2.50 0.57 4.37 1.72 0.38 1.77
Fold(MPM121/A veBPH) 1.52 0.74 29.40 28.99 19.31 22.88 2.31 16.83 0.41 2.77
veBPH) veBPH) 0.73 0.74 21.89 14.84 15.33 13.67 3.07 13.15 0.88 2.04
Fold(AveTumor/Av Fold(MPM120IA) Fold(
AveTumor 12.80 0.18 19.34 33.87 3.19 1.86 5.87 0.48 44.86
AveBPH 0.36 0.01 1.90 4.08 0.47 0.08 0.08 0.08 0.08
Secreted/Me mbrane Bound Maybe
Tissue Distribution spleen endothelial prostate breast colon bone colon
Gen Bank Accession Number N44111 AA158162 N48259 AA416627 T60111 H22932 R36066 N74131 AA005196 AA36142
IMAGE Clone ID 203579 500853 222080 731280 81331 52026 24815 268417 428086 75458

P O

Table 2-4: Poor vs. Good Clinical Outcome. Genes upregulated 3 fold or higher in any one tumor sample.

Table 24: Poor vs Good Clinical Outcome: Genes upregulated 3 fold or higher in any one tumor sample.

		Accession			Fold(AvePoor/A		Fold(mpm31/A Fold(mpm32/Av Fold(mpm33/A Fold(mpm34/A # or lumors Fold(mpm31/A Fold(mpm32/Av Fold(mpm33/A Fold(mpm31/A Fold(mpm31/A Fold(mpm32/Av Fold(mpm33/A Fold(mpm31/A Fold(mpm32/Av Fold(mpm33/A Fold(mpm33/A Fold(mpm32/Av Fold(mpm33/A Fold(mpm3/A Fold(mpm33/A Fold(mpm33/A Fold(mpm33/A Fold(mpm33/A Fold(mpm3/A Fold(m	Fold(mpm33/A veGood)	Fold(mpm34/A veGood)	# of lumors
Order	MAGE		AveGood	AVeroor	(poogev	(noone)				•
	Clone	Š				Č	57.56	34.98	23.80	,
•		04426470	1.42	42.16	29.71	70.7		7.7	3.64	က
_	0/050/		! !		80.8	0.69	10.04	•	7 73	es
8	124071	R02586	0.53	3.5		3.69	7.75	1.5.1	2 :	. «
C.	346321	W79647	9.0	3.29	5.17		6.82	0.22	6.80	o (
•	PEEROA	AA664101	24.18	108.55	4.49	7	10	4.66	0.57	m
* 1	10000	554073	13.03	52.25	4.01	4.90	- 6	1 30	3.29	က
S	42325	20102	3	6 33	2.97	4.00	3.29	3	9.60	m
ဖ	287528		2.13	9:35	267	0.45	3.61	3.01	9.0	
7	756595	AA444051	1.46	3.89	70.7	700	62.30	0.37	62.14	y (
. α	457374	AA700876	0.64	19.98	31.26	1 70 0	25.33	12.26	0.86	7
•	257105	W93315	0.87	8.54	9.81	10.0 10.0	47.80	3.83	0.33	7
D (A A 7004 E 2	1 24	6.95	5.61	0.47	90.5	14.0	10.17	7
9	460788	7002		4	5.49	0.82	10.19	5		2
F	284601	N76201	0.27	2.		0.02	8.87	0.03	6.63	. ‹
5	460487	AA677706	53.51	237.74	4.44	1 do 0	7.66	0.75	7.64	7
ī	240442	N58558	0.86	3.63	4.25	C:	4 6	2.47	6.75	8
2 :	21047	069949	55	2.37	4.23	0.93	27.0	, q	6.98	8
4	091717	10001		94	3.95	1.06	1.58	• (07.9	2
1 5	782513	AA432030	4 80.4	80. i	3 83	1.37	1.26	6.00	2.0	
9	782513	AA432030	2.03	7.78	3 6	0.92	3.71	0.70	3.87	4 (
17	129128	R10935	0.38	1.47	3.81	1 4	4.21	2.50	5.95	7 (
ξ.		AA630084	6.04 40.04	21.23	3.52		5.87	0.66	5.86	7
5 6		AA001431	1.47	5.16	3.51	7 6	464	0.26	6.62	74
2 6		AA677388	0.60	2.09	3.47	(c.)	. ע ע	0.00	4.97	7
3 6		N92842	0.05	0.16	3.28	60. F		0.36	5.81	8
. 6			1.20	3.83	3.19	0.73	786	1.37	4.54	8
4 6				12.41	3.17	CA:C	5 6	4.93	2.18	8
3 3		•		2.53	3.17	3.39	 	140	5.31	73
77	-	-		160 25	3.15	1.55	5.32	.	09.0	2
22	740941	•	4,7	5.5	3 04	1.25	4.71	5.51	80.0	, (
56	241705	H91680	1.71	5.18	r 6	2.54	3.88	4.35	1.25	4 (
27	754559	AA406301		. 96 90. 1	90.0	151	4.72	0.91	4.72	v (
28	3 487327	AA045524		1.46	06.2	1 27	2.66	3.34	4.49	7
i c	843098	AA488676	5.29	15.54	2.94	1				
í						-				

71	7	7	8	7	7	7	8	74	7	7	8	7	8	7	8	7	8	8	8	8	7	64	7	0	N	7	7	7	7	-	-	-	Ψ-	▼-	
4.79	4.42	4.95	5.26	4.59	0.00	3.66	3.81	2.47	3.92	4.19	3.45	3.21	3.86	0.87	1.24	3.83	3.59	4.17	3.58	3.02	3.39	3.49	3.19	3.32	3.63	3.16	3.44	3.20	3.19	1.47	1.63	15.38	0.82	0.31	
1.18	4.07	4.85	0.29	0.83	2.12	2.04	1.06	3.41	3.94	0.94	3.06	1.85	3.21	3.08	0.91	0.53	3.21	0.41	1.65	1.72	0.34	0.72	0.97	1.47	3.21	0.53	0.36	0.95	0.51	31.70	4.	2.08	0.64	0:30	
0 V	1.80	0.86	5.28	4.59	4.	3.67	3.82	1.28	1 43	4.20	1.25	3.22	1.22	4.30	3.72		1 07	4 18	9 G	3.03	3.34	3.49	3.16	3.33	0.69	3.16	3.45	3.22	3.20	0.65	1.63	0.71	1.02	17.43	
ç) 4) (1.52		4 43	- n) (1 (1 (1)	- -	37.1	0.50	0.92	, t		? ;	- - -	5 6	0. t	5 4	04.	0.5/	0.42		5.0	U.94	02.	8 C	- C	5 4	; ¢	2 6	, d	9. 9. 16. 87	10:01	0.00	10.11	2
	2.94	2.90	2.87	2.86	2.79	2.67	2.64	2.60	2.59	2.56	2.50	2.43	2.43	2.35	2.35	2.34	2.34	2.34	2.34	2.31	2.29	2.21	2.16	2.15	2.13	2.10	2.06	1.95	1.88	38.	8.67	5.39	4.69	4.65 6.00	4.63
	0.65	4.08	4.26	3.16	1.86	0.02	43.42	1.70	3.12	3.00	19.45	15.91	0.19	8.05	7.96	69.88	1.28	5.70	4.88	48.48	4.15	0.13	1.05	0.44	4.37	4.63	0.92	1.35	60.0	7.71	6.43	3.02	7.68	17.08	7.06
	0.22	1.41	1.48	1.10	0.67	0.01	16.45	0.65	1.20	1.17	7.79	6.54	0.08	3.43	3.39	29.86	0.55	2.44	2.09	20.98	1.81	0.06	0.49	0.21	2.05	2.20	0.45	0.70	0.05	4.15	0.74	0.56	1.64	3.68	1.53
	N73958	AA456886	AA432049	W46900	AA682386	AA436317	AA428415	W84716	AA701519	AA489743	AA485508	AA411384	R56898	AA598795	AA916325	AA011215	AA010221	AA041499	W42812	AA434502	W20458	R53558	R63646	H24350	H11063	AA452278	AA461174	H86554	AA053684	AA401441	AA455988	N90051	N51498	AA460511	H25546
•	298303	815542	782537	324437	461759	754393	770983	415692	435820	823696	811000	754600	41214	980868	1473304	359835	430231	376516	323238	770880	303089	39959	138788	52071	47418	787938	796806	223350	510136	741977	812074	305809	282104	795809	161456
	30	3.	32	33	8	35	36	37	38	38	4	4	42	43	4	45	46	47	48	49	20	5	25	23	2	22	26	22	28	29	8	6	62	63	8

	- •	-	-	-	-	-	-	-	-	-	······································	 .	,		·	-	-	-	-	-	-	₩.	-	▼	-	••	-	-	,- ·	• •	-	-	-	-	
12.08	0.42	1.00	14.50	13.33	13.47	0.62	1.33	11.93	1.09	1.48	13.64	0.61	2.41	8.86	0.48	0.84	0.20	9.83	0.73	8.89	1.16	0.79	0.45	1.30	1.20	1.27	0.51	6.89	1.7	1.55	1.10	2.15	2.30	2.18	
1.16	0.97	13.86	0.84	1.26	0.83	12.60	2.26	0.90	12.18	0.80	0.40	1.26	1.81	2.20	0.70	0.66	0.44	0.86	0.34	0.96	0.78	2.68	1.13	1.64	1.08	1.00	7.98	2.38	1.72	0.77	2.52	0.62	1.70	1.42	
1.61	0.92	0.70	1.15	76	06.0	0.52	10.78	1.21	1.09	1.48	0.31	11.31	1.97	1.14	12.18	0.92	96 0	0.92	0.73	2.00	80 80 10	1.70	1.47	1.30	1.20	1.81	0.57	06:0	7.08	2.28	2.89	2.16	2.01	1.41	
2.95	15.34	5 5	 	- 6	5.50	0.07	C	60.	0.62	0.02	0.43	5 5	7.07	1.51	2.0	0.40	10.97	11.6/	1.04	11.10	O. 0.	S 8	0.00	0.01		7.07) . c	200		0.00 7.4	2 6	4. n	9.00	87.2	·
4 45	2 :	4 . - 1	4.3/	4.25	4.23	3.97	3.97	3.80	3.78	3.74	3.73	3.69	3.55	3.53	3.48	3.44	3.35	3.32	3.31	3.24	3.18	3.16	3.01	2.97	2.96	2.96	2.84	2.81	2.77	2.77	2.76	2.75	2.70	2.70	2.3
000	0.02	0.39	3.58	2.10	4.89	4.59	6.59	2.51	3.06	3.20	5.85	18.28	0.32	73.02	3.01	0.21	0.73	0.15	2.94	8.46	0.85	1.53	48.46	6.88	13.27	2.40	0.93	2.99	3.09	4.88	0.09	6.25	32.44	2.47	2.89
!	0.18	60.0	0.82	0.49	1.16	1.16	1.66	99.0	0.81	0.86	1.57	4.95	60.0	20.71	0.87	90.0	0.22	0.04	0.89	2.61	0.27	0.48	16.11	2.32	4.48	0.81	0.33	1.08	1.1	1.76	0.03	2.28	12.01	0.91	1.07
	AA782380	AA863204	R63823	AA704718	W69960	H41222	H19128	AA628410	N48261	AA425382	W38679	AA451886	H06093	N64840	T77410	W47487	N33041	T99719	AA456413	W48762	H38425	H12981	AA136808	AA878880	N91797	N47738	H17882	H44861	W31255	R00480	AA160606	AA133721	AA486281	AA598659	AA024898
	857442	1455740	141336	450948	343919	192258	50930	1032774	282117	773157	320857	786657	44351	284701	23790	324313	270343	122782	788386	325128	192401	43207	491001	1493160	307174	277423	50182	176904	306540	123331	592707	512116	842848	106/68	365227
	65	99	67	. 60	0 0	2 8	7	72	73	7.	75	76	11	78	26	2 8	8 2	5 &	3 8	3 2	35	9 6	87	8	8	8	91	95	69	94	95	96	97	86	66

Table 2-4: Poor vs. Good Clinical Outcome: Genes upregulated 3 fold or higher in any one tumor sample.

-	-	-	-	-	-	-	-	-	-	-	-	-		- -	-	-	~ -	-	-	-	-	-	-	-	-	-		-	-	-	-	-	-	-	
6.67	0.71	1.74	6.57	1.96	0.94	1.75	1.41	1.54	2.93	2.01	1.15	2.36	1.80	6.80	3.38	2.70	0.26	0.55	1.81	1.33	1.39	1.49	2.10	0.72	0.69	1.24	6.39	1.59	6.65	2.40	1.29	0.89	1.26	0.54	
96.0	1.12	1.48	0.98	2.34	0.92	2.75	1.08	1.45	4.64	1.56	0.77	1.72	0.69	1.41	2.86	2.51	0.50	1.03	4.85	1.45	3.61	2.19	2.30	2.03	1.21	1.94	0.72	1.60	0.98	1.88	4.92	1.83	0.53	1.25	
1.47	1.13	790	44	1.14	53	22.1	2.25	2.48	1 12	7.70	1.48	2.39	1.81	0.93	2.51	3.23	0.86	1.47	1 90	2 2 2	2.05	3.76		20.5	1.32	2.52	1.00	1.11	. ¥	0.52	1 29	9 7	0.65	141	<u>:</u>
4 R7	5. t	7.70	0.70	7.87	9 6	n (0	4.00 4.00		4.00	. 4 6 6, 7			1 &	2 6	7 7 6	 	0.00	0.20	4 66	07:1	D 00	00.7	2.20	0 0 0 0	0.90		0.00	 		16.0	 	00.0	0 9 0 9	n 60 0 0	0.00
ć	2.69	2.68	2.67	2.66	2.60	2.60	2.56	2.55	2.54	2.54	2.52	2.50	2.50	2.49	2.48	2.47	2.45	2.45	2.45	2.44	2.44	2.43	2.43	2.41	2.39	2.39	2.3/	2.3/	2.36	2.35	2.34	2.34	2.34	2.33	2.32
	2.38	2.19	2.51	3.75	21.09	0.73	4.25	7.65	3.12	7.59	14.37	13.42	0.07	34.17	1.57	2.60	5.92	23.38	6.48	7.65	0.13	2.62	3.96	6.17	1.65	4.56	4.66	2.42	0.38	1.74	30.32	26.12	0.54	0.13	6.68
	0.88	0.82	0.9 46.0	1.41	8.12	0.28	1.66	3.00	1.23	2.99	5.71	5.37	0.03	13.72	0.63	1.05	2.41	9.53	2.65	3.13	0.05	1.08	1.63	2.56	0.69	- 1.91	1.96	1.02	0.16	0.74	12.93	11.15	0.23	90.0	2.88
	AA701931	AA885311	H29256	N51367	N79738	H75860	AA682631	R17962	W76603	R92601		AA010932	AA676470	AA443004	R15409	T95274	AA865464	AA621031	AA629991	AA434362	AA704844	AA278408	AA778098	AA421389	AA999953	N27992	N76101	AA628146	R12183	AA757732	R98064	R11587	AA621324	W02342	N26163
	435567	1461664	49630	283208	289599	230560	431296	32381	345670	196257	361363	359781	882511	811766	53162	120162	1470048	1056203	884514	837908	453218	703930	379920	739237	1636495	265102	299412	1055737	25590	396111	206838	130046	1048794	292071	269411
	100	101	102	103	5	105	90	107	108	109	19	77	112	113	1 4	115	116	117	118	119	120	121	122	123	124	125	126	127	128	129	130	131	132	133	134

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1.28	0.83	0.78	2.08	0.52	0.76	0.43	185	35.		04.1	1.35	3.42	1.52	0.51	0.80	1.86	1.66	0.81	1.55	0.53	3.05	5.49	1.70	5.74	1.18	1.09	1.09	4.10	2.24	1.13	0.95	2.18	0.92	3.17	0.97	
2.50	1.79	1.52	2.27	1.21	5.15	. t		31 1	1.45	2.79	1.20	1.63	2.50	0.95	0.8 4	1.71	3.14	1.03	<u>;</u>	4.	2.50	1.15	1.33	40.1	<u>4</u> .	1.52	2.08	1.37	0.92	0.92	0.90	<u>4</u> .	5.68	1.66	5.76	
0.87	2.66	1.74	3.23	162	2		7.77	08.F	0.75	3.29	1.80	2.82	1.36	1.13	6.45	1.57	1.13	1.02	1.67	1.92	2.59	0.86	0.67	1.13	1.55	2.34	2.19	1.13	2.24	1.13	1.29	1.52	1.29	1.19	0.97	
4.59	3.93	5.17	62.1) (a)	3 6	70.7	BS:4	2.29	6,15	1.56	4.74	1.21	3.67	6.43	0.88	3.81	2.99	6.07	4.18	4.93	0.68	1 28	5.06	0.80	4.62	3.71	3.29	2.04	3.19	5.41	5.44	3.03	0.67	2.51	0.82	
2.31	2.30	2.30	0000	67.7	67.7	87.7	2.29	2.28	2.28	2.28	2.27	2.27	. 2.26	2.26	2.24	2.24	2.23	2.23	2.21	2.21	2.20	2.30	2 19	: c	2.17	2.17	2.16	2.16	2.15	2.14	2.14	2.14	2.14	2.13	2.13	ų
8.92	23.49	52.52 87.4	t (*)	4 (A	0.13	4.17	19.01	1.33	0.75	4.37	4.23	1.08	0.98	1.52	76.0	2.12	3.14	0.76	0.05	8 5	3.38	3	 	50.0	2.03 7.96	1.80	28.92	5.24	8.31	60.65	1.22	5.69	0.58	11.18	1.14	
4 29	600	22.01	0.70	3.90	90.0	1.82	8.32	0.58	0.33	1.92	1.86	0.48	0.43	0.68	0.42	980	4 44	7, 6	5 6	7 O	6. 4 4. 4	3 6	0 0	9 6	20.0 40.0	3 8	13.37	2.43	3.87	28.28	0.57	2.66	0.27	5.24	95.0	
A A 708201	00107344	AA197344	AA873089	W76331	AA444111	AA676899	AA011593	H15250	R44776	AA733090	N45398	AA406059	N95435	H25689	N55430	A A 598659	A A 677206	AA621467	A A C 4 7 8 4 8	AA024844	1401/640	Rosoad	AA2/9409	AK142913	AA004073	NEGRIF S	AA171899	AA134862	AA454165	AA011095	AA419612	H53141	AA777417	AA251386	AA670429	· · · · · · · · · · · · · · · · · · ·
202708	034400	999/79	1325/521	345055	757135	455179	429678	49687	33096	399577	260187	743026	30000	462077	246936	807004	00,100	404/30	276020	2/00/2	1564505	180023	704227	504420	571405	7076R7	610374	502436	795322	359661	746148	202521	449061	684562	878836	
104	2 4	2 5	13/	138	139	140	141	142	143	144	145	146	147	α 7	2 2	1 4	0 4		707	3 4	\$;	60	136	12/	200	2 2	5 6		7 2	164	20.00	8 6	167	2 4	2 2	<u>}</u>

Table 2-4: Poor vs. Good Clinical Outcome: Genes upregulated 3 fold or higher in any one tumor sample.

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99.	4.82	0.84	1.32	1.89	4.53	1.65	3.11	2.79	1.39	0.25	3.92	1.44	1.78	0.62	5.97	0.15	1.50	1.36	0.84	1.12	1.74	1.34	2.51	0.94	0.44	1.67	0.59	0.69	2.01	0.65	0.65	0.68	1.39	0.62	
00	69.	1.07	4.86	0.92	1.88	1.47	1.24	3.06	1.24	2.62	1.57	0.96	3.50	1.09	0.71	0.13	1.68	3.03	0.70	2.17	1.67	1.57	0.12	3.10	0.99	1.23	1.34	1.49	0.99	1.52	1.72	1.53	4.20	1.31	
,	1.33) iii	58.4	5 5	2 2	1.65	1.16	36	3.23	0.86	0.73	1.79	50.1	0 60	7.72		67:	30 5.44	t ;	47.4	4.37	, 75 10 10 10 10 10 10 10 10 10 10 10 10 10	2.82	2.51	4.23	1.63	5 6	5 6	1 97	10.1	, i	1 19	1.22	
	1.07	1.07	07.0	1.60	3.70	ж Э	9 T. C	, , ,	 	0.4 0	£.13	t		0 1	7.60	0.30	0.20	3.72	2.42	1.18	3.74 4 4	3.01	0.0 4 4 %	0.10	1.27) a	0.30	6. 4. 6. 6. 6. 6. 6. 6. 6. 6. 6. 6. 6. 6. 6.	- c	8.00 8.00 8.00	9 t	, i.o.	5.7g	1.20	} •
	2.12	2.12	2.11	2.11	2.10	2.10	2.09	2.08	2.08	2.07	2.07	2.07	2.07	2.06	2.06	2.05	2.05	2.05	2.04	2.04	2.04	2.04	2.03	2.03	2.03	2.03	2.03	2.03	2.02	2.02	2.02	2.02	2.02	2.02	Z.U1
	0.78	90.65	3.29	13.63	13.41	3.21	2.34	27.88	2.66	0.79	36.21	0.69	1.92	3.48	2.14	16.88	9.61	6.73	1.09	0.27	2.06	3.34	0.75	18.02	13.69	4.76	0.77	5.26	4.26	1.38	26.52	9.77	7.47	3.17	0.24
	0.37	42.83	1.56	6.45	6.39	1.53	1.12	13.37	1.28	0.38	17.47	0.33	0.93	1.69	. 2	8.23	4.69	3.29	0.53	0.13	<u>.</u>	2 .	0.37	8.88	6.75	2.35	0.38	2.60	2.11	0.68	13.11	4.84	3.71	1.57	0.12
	AA706955	W96187	H83283	AA626242	AA010223	AA521414	N47308	AA455565	AA453040	AA676948	AA876375	N71769	AA187789	AA030046	W58013	R45964	AA031513	AA011185	AA430242	96660H	T61792	AA676840	AA872602	AA151002	AA968896	N20328	AA598572	AA609983	AA164782	AA150507	W48700	H67292	AA490985	R10948	H09749
	451898	361668	199175	745524	430235	826099	280508	813488	788574	460189	1493107	290702	624379.	469969	341336	35575	470393	359713	781091	46916	78946	460114	1323328	504927	1574594	264597	897822	1031976	595001	491763	324885	229573	824588	129148	46453
	170	12	172	173	174	175	176	177	178	179	180	181	182	183	18 48	185	186	187	88	189	190	19	192	193	194	195	196	197	198	199	200	201	202	203	204

Table 2-4: Poor vs. Good Clinical Outcome! Genes upregulated 3 fold or higher in any one tumor sample.

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75		47.1	1.02	0.90	0.79	0.59	3.62	1.46	1.1	0.80	3.20	1.30	0.75	0.51	1.50	1.78	1.31	0.73	1.30	6.57	2.06	0.82	1.01	0.97	0.93	3.07	1.09	0.89	1.10	0.93	1.65	1.13	0.97	2.39	60.0	
4 25	5 6	7.50	1.20	1.69	1.29	1.17	1.20	4.84	3.49	1.22	1.25	1.07	1.82	1.38	1.09	1.28	3.99	2.79	3.96	0.42	2.03	1.05	0.93	1.03	1.19	2.59	1.15	0.76	1.00	1.26	3.06	4.27	1.19	3.27	0.67	
7	C/:1	0.76	96.0	1.88	4.31	5.08	1.38	0.80	2.01	1.69	4.1	1.55	0.76	0.95	1.33	42.	1.11	3.06	1.63	0.40	3.64	3.99	1.02	0.98	0.94	0.82	1.10	2.06	2.00	1.74	1.80	1.14	1.77	1.18	6.77	
ć	3.60	3.49	4.80	3.49	1.56	1.09	1.71	0.81	1.29	4.19	2.02	3.98	4.53	5.03	3.92	3.22	04.1	1.22	0.87	0.36	0.00	1.86	4.75	4.71	4.62	1.20	4.33	3.95	3.55	3.70	1.10	1.07	3.66	0.75	0.05	
1	2.00	2.00	1.99	1.99	1.99	1.98	1 98	198	80	9 6 6	86	1.97	1 97	1.97	1.96	1.96	195	1.95	1 94	, t	1.93	1.93	1.93	1.92	1.92	1.92	1.92	1.91	1.91	1.91	1,90	1.90	1.90	06.1	180	^
	5.87	9.35	0.91	19 17	2.63	0.16	27.07	2 2 6	5 5	i	9. E	0.25	200	600	2.75	7.4		33.80	200 F	5.50	0.02	4.71	0.54	1.01	0.41	4.14	0.68	1.96	55.28	6.25	1.81	0.56	46.8	3.29	2 5) ř
	2.93	4.68	0.45	680	5 6	200	3 4		? · · ·	4.40 4.40	4. 4 0. 4 0. 4	3 5	2 2	\$ 2	1.60) i	0.0	47.37	, i	20.0	2 5	2,5	20	53.0	0.21	2.16	0.35	1.03	28.91	3.28	0.95	0.29	4 7.1	473		<u>.</u>
	T83657	N51362	R31114	AA195801	007048	M02073	20204	1425000 A A B 7043E	A 6 6 4 0 6 4 0 6 4 0 6 4 0 6 4 0 6 4 0 6 6 4 0 6 6 4 0 6 6 6 4 0 6 6 6 6	A4664103	WISSIS	0 0 4 5 5 8 2 5		14732730	AA250115	A A 450473	CA169175	N36263	* COSCO *	AA588455	H04828	A A 634379	772007	H29290	R39364	AA449048	H20717	R01245	AA453105	A A 478606	H03146	9067W	R92801	R23609	174064	
	113284	283191	134265	CER407	121000	227262	747 770	/50007	1505060	80000	322643	644600	2010	277405	CBER64	0000	90000	120062	128833	720099	430EA	743828	22240	40858	24632	785816	51420	124137	789091	75363	181897	346968	197056	132140	07070	2
	205	206	207	2 0	000	2 6	2 7	- 6	717	513	214	246	1 5	240	240	9 6	2 6	ן קל נל	777	3 5	724 225	228	227	328	220	230	231	232	233	23.5	23.5	236	237	220	200	200

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3.84		1.20	3.33	0.61	0.86	1.70	1.01	1.21	0.89	3.29	1.22	1.05	2.90	5.28	3.29	1.52	0.63	3.13	1.14	0.96	0.66	1.47	3.60	0.11	0.63	4.1	1.33	0.71	1.30	1.21	4.80	1.39	0.92	3.23	3.61		
4 78	8 .	1.19	1.50	1.46	4.52	4.05	1.38	144	3.79	1.70	1.07	44.	0.16	0.85	1.62	3.89	28.	2.00	1.22	1	1.72	61.1	0.73	0.82	1.26	1.06	1.27	1.17	3.76	3.39	0.90	3.02	1.79	4	1.09	}	
6	1.03	1.68	1.70	2.06	60 -) (4)	90.0	- 1 0 %	-:00		123	901	0.4	0.73	146	0.71	162	20:- 28:-		S 4	2 7	÷ 5	1 28	. c	1.07	690	1 27	. 6. . 6.	1.22	127	180	. 8	0.50	26.0	7.3 2	2	
	0.85	3.48	101	000	5 5	<u> </u>	07.1	3.00	0.30 0.00	S 10	7.0	, c	9/.0	0.20	5 6	0.97	7.7	3.10	1.30	3.23	3.64	3.35 3.35	3.20		5.0 F	÷ •	3 6	3.22	(a c) T	- c	G 6	1.26	14.5	04.1		œ
	1.89	1.89	9 8	8	1.88	1.88	1.88	1.85	1.85	1.85	1.84	48.	1.83	1.83	1.83	.83	1.82	1.82	1.82	1.81	1.79	1.79	1.79	1.78	1.78	1.78	1.78	1.7.	<u> </u>).'L	1.77	1.77	1.76	1.76	1.75		w
	2.11		D (0.89	53.13	2.13	2.63	3.49	8.26	1.09	1.38	41.96	0.53	5.23	1.90	1.56	2.99	2.48	2.68	12.19	2.78	2.18	2.31	2.53	0.03	2.80	8.55	0.87	4.09	7.89	2.64	5.15	0.26	7.01	2.78	10.90	
	1 12	- · ·	3.97	0.47	28.27	1.14	1.40	1.88	4.47	0.59	0.75	22.82	0.29	2.86	<u>5</u>	0.86	1 .6	1.36	1.48	6.74	1.55	1.22	1.29	1.42	0.01	1.57	4.81	0.49	2.31	4.46	1.49	2.92	0.14	3.99	1.59	6.25	
	T64854	100101	AA453531	AA682321	AA443698	AA453863	R98344	AA155787	AA476961	N20989	AA704802	AA664406	AA609982	R38645	AA772497	AA677984	N91744	AA282594	AA434117	AA677397	W88623	AA620421	AA48194	N46759	AA885052	R71124	AA030004	H61082	AA620611	R53342	AA905838	AA827405	AA126682	R70684	AA876354	W05033	
		78680	788136	462926	784016	813719	201641	590253	739578	265853	452468	868652	1031203	23000	413300	430763	293328	713019	770216	454896	417691	951022	782797	277112	1466485	142944	470216	206172	1048702	40031	1505294	1422794	502593	141815	1493079	298648	
	6	240	241	242	243	244	245	246	247	248	249	250	251	252	253	254	255	256	257	258	259	260	261	262	263	264	265	266	267	268	269	270	271	272	273	274	

Table 2-4: Poor vs. Good Clinical Outcome. Genes upregulated 3 fold or higher in any one tumor sample.

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0.89	1.36	1.28	0.48	1.07	3.13	0.81	0.28	3.32	1.21	0.40	0.71	3.71	0.73	4.33	3.77	3.29	0.53	3.62	0.84	3.67	0.83	1.05	0.25	0.45	1.25	3.00	1.76	0.16	3.43	1.35	1.17	0.72	0.99	1.24
1.11	0.81	0.93	1.10	1.53	1.38	1.81	0.87	1.21	3.37	1.42	1.42	1.47	1.70	1.21	1.07	1.41	4.89	1.33	1.09	1.42	0.95	0.99	0.76	1.78	1.12	1.46	0.62	0.91	1.33	1.31	1.15	1.65	1.23	1.01
1.24	1.36	1.28	2.26	0.59	1.60	3.25	2.31	1.20	1.22	1.45	3.33	0.78	1.43	0.78	1.18	0.65	0.51	0.95	1.63	1.03	0.57	1.50	4.85	3.15	1.26	1.11	0.87	2.51	1.01	0.81	1.18	1.21	0.80	1.1
3.73	3.43	3.48	3.11	3.75	0.81	1.05	3.45	1.16	1.08	3.62	1,41	0.91	3.01	0.53	0.83	1.50	0.91	0.94	3.27	0.70	4.47	3.29	0.95	1.42	3.17	1.22	3.52	3.15	0.95	3.23	3.18	3.11	3.66	3.31
1.74	1.74	1.74	1.74	1.74	1.73	1.73	1.73	1.72	1.72	1.72	1.72	1.72	1.72	1.71	1.71	1.71	1.71	1.71	1.71	1.71	1.70	1.70	1.70	1.70	1.70	1.70	1.69	1.68	1.68	1.68	1.67	1.67	1.67	1.67
7.07	33	4.53	3.02	29 07	2.13	57.00	5.19	1.38	0.58	35.33	5.05	1.42	7.12	5.74	0.69	2.60	0.82	1.74	62.95	2.09	0.33	21.50	5.98	3.24	1.78	3.16	0.06	0.07	0.90	10.94	47.59	5.91	8.82	0.85
30	2 2	5 6	77	18.74	1.23	32.95	3.00	0.80	934	20.52	2.94	0.82	4.15	3.35	0.40	1.52	0.48	1.01	36.82	1.23	0.19	12.62	3.51	1.91	1.05	1.86	0.03	9	45.0	6.53	28.46	8. 40.	5.28	0.51
4447344	T81289	44457235	AA057032	P09179	AA455980	AA62095	R38943	W71990	N26608	AA459761	AA255551	H80063	W73724	R81831	AA700843	AA701948		AA464598	AA047260	W86124	R54179	AA456695	A1003028	H96392	H94563	R82317	H84734	H73806	AA437093	AA464068	AA664389	AA733061	AA420992	AA883114
724469	26764	10007	400043	427640	812053	1056172	24958	345701	269172	796366	682477	249070	344108	147834	436097	435919	25122	812961	488964	416227	41608	813149	1684274	256680	243159	149058	220135	214713	757325	810284	868630	399562	731255	1466942
376	276	27.0	270	270	280	2 2 2 2	2 6	283	284	285	286	287	288	289	290	291	200	203	294	295	950	207	298	292	300	30.	302	303	30,	305	306	307	308	309

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3.30	1.22	0.71	3.88	1.61	1.38	1.38	0.87	1.1	0.62	3.05	1.68	0.34	0.68	1.58	0.84	5.54	0.94	1.18	0.83	1.11	0.85	0.65	0.80	3.88	0.91	0.48	0.84	0.70	0.79	1.24	0.79	0.55	1.32	1.19	
1.25	1.08	1.09	1.29	1.06	1.08	3.11	1.45	1.06	1.32	1.14	1.21	0.77	0.75	1.24	1.03	0.28	0.82	1.19	0.58	1.25	4.10	1.23	96.0	0.91	3.56	0.88	1.23	1.60	40.1	0.57	0.88	0.41	1.31	0.91	
0.92	1.21	1.00	990	0.61	88.0	70.4	300	1.20	1.40	1.57	0.58	2.19	1.37	0.59	0.84	25.0	1 20	3.28	1.15	96.0	72.0	1.45	40.	0.86	0.84	1.61	1.25	3.05	1.38	1.24	0.81	4.59	0.58	0.86	i
1 20	 		8 6	0.00 36		3.50	 5	 	5.25	5.23 2.48			5.66 7.76	, c		9 9	9 6	3.37 80	0.00	5.00 5.41		- 80 - 60 - 60				. or r	3 03	8 6		3. 15 3. 25	2 6	0.00	t 70	40°6	
ì	7.67	1.6/	1.66	1.66	1.66	1.65	1.65	1.64	1.64	1.64	49. 1.64	1.63	1.63	1.63	1.62	1.62	1.62	1.62	1.61	1.61	1.61	1.61	1.60	9.50	8c. 1	60. T	90.T	6	3.58	1.58	2.58 86.1	1.58	1.57	1.56	1.35
	0.23	0.57	0.15	42.64	2.46	2.82	0.59	4.02	0.80	7.92	0.71	0.28	12.12	0.76	9.47	5.76	12.39	17.82	0.89	3.93	3.25	2.10	4.80	0.33	5.92	3.22	2.95	26.88	2.27	7.70	2.42	11.90	1.36	1.75	1.76
	0.14	0.34	0.09	25.66	1.48	1.70	0.35	2.44	0.49	4.83	0.43	0.17	7.43	0.46	5.83	3.54	7.64	11.02	0.55	2.45	2.02	1.31	3.00	0.21	3.72	2.03	1.86	16.93	1.43	4.86	1.54	7.55	0.86	1.12	1.14
	AA884397	AA788738	AA983362	AA757417	T65398	R89358	H48138	H16439	AA719257	AA479512	AA279990	H15040	AA504239	H68655	R43553	AA150422	R43053	AA012867	AA459472	AA774665	W73748	AA487676	AA070435	N48913	AA156926	AA406019	AA453900	AA116061	R37357	AA169606	H11036	AA279145	AA456377	H24580	H23548
	1461725	1240347	1592715	395417	21634	196038	193780	47527	1292593	754313	705064	48454	825031	212325	32794	491415	32509	360885	814460	856289	344272	841624	530954	279613	502518	742132	813755	531862	28270	594266	47378	703943	813256	160730	52002
	310		312	313	314	3,15	316	317	318	319	320	321	322	323	324	325	326	327	328	329	330	331	332	333	334	335	336	337	338	339	340	34.	342	343	344

-	-	-	-	-	-		-	-	-	-	•	-	-	-	•	-	-	-	-	-	-	-		-	-	-	-	-	-	-	-	-	₩.	-		
0.81	3.45	0.47	3.94	3.79	1.26	0.42	0.72	1.26	0.83	0.70	4.14	3.11	3.79	0.85	0.43	3.36	0.22	3.08	0.78	0.80	3.69	3.76	0.31	0.61	0.86	0.69	3.22	0.76	0.56	0.78	0.50	0.77	0.58	0.37	· •	
0.93	1.06	0.66	0.86	0.68	3.21	1.98	0.75	306	76.0	1.48	0.56	1,13	0.71	3.38	1.63	0.74	0.76	0.95	0.80	0.67		220	0.52	1.12	0.91	41.1	0.96	0.81	0.59	0.60	0.57	0.71	0.48	0.74	Š	
1.16	0.90	1.45	0.70	9.:0 0%.0	5 6	. 4 . 4	2 8		3. t		250	5 5	28.0) c		9 6	8	+-7	9 6	D (C	5.03 25	5 9	00 T	47.0	 	5.75	0.50	0.79	0.87	1.19		S 5	S: •	5.00	7.1	
3 28	78	0 0	9.00 0.00 0.00 0.00	6.50 6.00 6.00	90.6	n (6 5 (9.0 6.0	50.50 10.00	0.0	9. O	9 6	0.03	0.0	60.0	0.83	0.70	0.68	8. 8. 8. 8. 8. 8. 8. 8. 8. 8. 8. 8. 8. 8	Q (3.45	3.31	0.86	0.35	3.35	3.20	0.7°	. S	- e	9 6	9 6	† 0	0.88 80.00	3.08 1.08	3.47	3.29	
i i	6	1.55	4. j	2	1.54	1.54	1.54	1.53	1.52	1.52	1.52	1.51	1.51	1.50	1.50	1.47	1.47	1.46	1.46	1.45	1.45	1.44	1.43	1.43	1.43	1,43	1.43	74.1	14.1	04.1	1.40	1.40	1.40	1.40	1.39	÷
:	2.13	1.22	9.93	3.15	2.24	0.16	1.41	8.37	0.61	2.19	1.42	5.60	2.09	1.58	1.47	0.09	0.73	14.17	1.51	0.16	0.70	1.98	3.43	11.76	2.63	0.09	0.1	4.22	1.23	0.63	16.09	1.11	12.76	0.80	1.83	
	1.38	0.79	6.43	2.04	1.45	0.11	0.92	5.46	0.40	1.	9.0	3.70	1.39	1.05	0.98	90.0	0.50	9.68	<u>+</u> .	0.11	0.48	1.37	2.40	8.21	4 .	90.0	0.08	2.96	0.87	0.45	11.47	0.80	9.13	0.57	1.31	
	H59188	W24646	AA282159	AA620466	AA400002	T69522	R16541	AA682274	AA779225	R93124	AA886792	AA406320	AA278842	AA912071	N50904	AA161248	H51122	R98472	AA411668	H10073	AA779212	R73088	H02294	H99672	AA644563	AA004671	R01937	H06377	AA677025	R92994	AA193254	W86183	AA428014	N54788	R89082	
	204251	308223	712888	951108	743187	66952	129589	461699	452676	196992	1501546	754591	703827	1486752	281045	505414	179631	201125	753271	46844	453988	141548	151184	263697	845521	428796	124567	44300	454564	196612	665774	416305	773483	244310	195751	
	345	346	347	348	349	350	351	352	353	354	355	356	357	358	359	360	361	362	363	364	365	366	367	368	369	370	371	372	373	374	375	376	377	378	379	

0.99 3.23 3.78 0.31 0.76 0.51 0.49 0.49 0.40 0.40 0.40 0.40
0.62 0.69 0.73 3.28 0.73 0.70 0.73 0.59 0.64 4.01 0.64 0.36 0.36 0.36
3.15 0.80 0.37 1.04 0.78 3.48 3.86 0.70 0.53 0.45 0.45 3.65 3.40
0.53 0.74 0.78 3.35 0.34 0.59 3.50 3.11 3.12 0.13 3.29 0.13
1.38 1.36 1.36 1.35 1.35 1.32 1.32 1.29 1.29 1.25 1.20 1.09
3.59 1.84 2.34 14.94 4.30 1.52 3.71 1.41 0.57 1.30 5.27 9.28 2.54 7.36 0.10 6.03 2.20 4.23 2.5.11
2.61 1.35 1.72 11.02 3.20 1.13 2.79 1.07 0.43 1.01 4.08 7.18 2.03 5.89 0.08 4.98 1.82 3.53 23.01
AA922309 AA706969 W93483 AA598815 T73883 N49629 AA102670 AA040265 AA478542 AA478542 AA443706 H49519 AA401137 AA6843592 AA6843592 AA465495
1474284 451911 357264 897952 84464 277708 243741 563598 376043 784772 222157 224100 42373 784017 178860 741497 380890 1390584 462325
380 381 382 382 383 384 385 390 391 392 393 393 396 396 398 398 398 399

Table 2-5: Poor vs. Good Clinical Outcome: Genes upregulated and or more in at least 2 tumor samples

Tabl4 2-5: Poor vs Good Clinical Outcome: Genes upregulated 3 fold or more in at least 2 tumor samples.

Fold(mpm33/A Fold(mpm34/Av # of Tumor veGood) Samples	•	ינים	က	က	က	က	က	ო	~	· ~			۰ ،	۱ ۵		. ~		4 (4 6	, ,	4 (4 (4 6	1 c	v (4 (7 (ν (N (7 (4 (:
Fold(mpm34 eGood)		23.80	3.64	7.73	6.80	0.57	3.29	3.60	62 14	, ag	9 6	6.33	2.0 40 40 40	7 6	2 4	2 6	9 9	9.70	78.5 18.5 18.5		0 0	20.0	, 4. v	0.0	40.4	2.18	5.31	0.69	1.25	4.72	9.4 .	70
Fold(mpm33/A		34.98	3.34	1.51	0.22	4.66	130	- «	9 6	0.37	12.26	3.83	0.77		0.70	2.47	81.0 1	6.00	0.70	2.50	0.66	0.26	0.00	0.36	1.37	4.93	0.41	5.51	4.35	0.91	3.34	•
Fold(mpm32/ AveGood)	(1)	57.56	16.64	7.75	6.82	19.5		87.5	3.61	62.30	25.33	17.80	10.19	8.87	7.66	6.76	1.58	1.26	3.71	4.21	5.87	6.64	6.55	5.82	0.84	2.18	5.32	4.71	3.88	4.72	2.66	
Fold(AvePoor/ Fold(mpm31/Av Fold(mpm32/	(noone	2.52	0.69	3.69	4 42	i 6	06.4	4.00	0.45	0.24	0.81	0.47	0.82	0.02	0.95	0.93	1.06	1.37	0.92	1.40	1.66	0.37	1.59	0.75	5.95	3.39	1.55	1.25	2.54	1.51	1.27	
Fold(AvePoor/	AveGood)	29.71	808	, u	- ·	D 1	4.01	2.97	2.67	31.26	9.81	5.61	5.49	4.44	4.25	4.23	3.95	3.83	3.81	3.52	3.51	3.47	3.28	3.19	3.17	3.17	3.15	3.04	3.00	2.96	2.94	
AvePoor		42.16	2 24	3.6	5.29	108.55	52.25	6.32	3.89	19.98	8.54	6.95	1.50	237.74	3.63	2.37	16.09	7.78	1.47	21.23	5,16	2.09	0.16	3.83	12.41	2.53	169.35	5.19	3.96	1.46	15.54	
AveGood		•	4 · ·	0.53	0.64	24.18	13.03	2.13	1.46	0.64	0.87	1.24	0.27	53.51	0.86	0.56	4.08	2.03	0.38	8. 29.	1.47	0.60	0.05	1.20	3.91	0.80	53.79	1.71	1.32	0.49	5.29	
Accession	No.	04700744	AA4504/8	K02586	W79647	AA664101	R61073		AA444051	AA700876	W93315	AA708152	N76201	AA677706	N58558	H68848	AA432030	AA432030	R10935	AA630084	AA001431	AA677388	N92842	W84751	AA099262	AA028171	AA478298	H91680	AA406301	AA045524	AA488676	
IMAGE	Clone ID		9/056/	124071	346321	855624	42325	287528	756595	452374	357105	460798	284601	460487	248412	212188	782513	782513	129128	854678	362059	454822	308696	415766	489535	364865	740941	241705	754559	487327	843098))
1 200		•	- (7	က	4	S.	9	_	· 00	σ	, C	=======================================	12	5	4	5	<u> </u>	2 5	<u>6</u>	6	2 2	21	2	3 2	5 2	י גי	3 %	27	; e	i s	7

0 0	7	8	2	7	74	7	7	74	7	7	7	63	7	2	2	8	7	7	8	8	7	7	7	2	2	7	2	
4.42	5.26	4.59	0.00	3.66	3.81	2.47	3.92	4.19	3.45	3.21	3.86	0.87	1.24	3.83	3.59	4.17	3.58	3.02	3.39	3.49	3.19	3.32	3.63	3.16	3.44	3.20	3.19	
4.07	0.29	0.83	2.7	2.04	1.06	3.41	3.94	0.94	3.06	1.85	3.21	3.08	16.0	53	3.24	0.41	1.65	1.72	98.0	0.72	0.97	1.47	3.21	0.53	95.0 36	0.95	0.51	;
1.80	5.28	4.59	3.14	3.67	9.82	1.08	143	4.20	1.25	22.5	22	4.30	3 5	4 00	3.00		2 G	8 8 8 8	 	3.49	9.50	88.6	9000	9 6	3.10 2.46		77.5	3.50
1.32	5 6	4.43	 5 .		-	27.1	3.20		8 6	. 4 8 6		= ;	1.13	94.5	1.13	1.48	0.57	0.42	04.1	0.70	. c.	9 ° °	6.09	98.0	1.38	4.0	0.16 0.16	0.53
2.90	2.87	2.86	2.79	2.67	2.64	2.60	2.59	2.50	2.50	2.43	2.43	2.35	2.35	2.34	2.34	2.34	2.34	2.31	2.29	2.21	2.16	2.15	2.13	2.10	2.06	1.93	1.88	1.86
4.08	4.26	3.16	1.86	0.02	43.42	1.70	3.12	3.00	19.45	15.91	0.19	8.05	7.96	69.88	1.28	5.70	4.88	48.48	4.15	0.13	1.05	0.44	4.37	4.63	0.92	1.35	60.0	7.71
1.41	1.48	1.10	0.67	0.01	16.45	0.65	1.20	1.17	7.79	6.54	0.08	3.43	3.39	29.86	0.55	2.44	2.09	20.98	1.81	90.0	0.49	0.21	2.05	2.20	0.45	0.70	0.05	4.15
AA456886	AA432049	W46900	AA682386	AA436317	AA428415	W84716	AA701519	AA489743	AA485508	AA411384	R56898	AA598795	AA916325	AA011215	AA010221	AA041499	W42812	AA434502	W20458	R53558	R63646	H24350	H11063	AA452278	AA461174	H86554	AA053684	AA401441
815542	782537	324437	461759	754393	770983	415692	435820	823696	811000	754600	41214	898096	1473304	359835	430231	376516	323238	770880	303099	39959	138788	52071	47418	787938	796806	223350	510136	741977
31	32	33	8	35	36	37	38	39	6	41	42	43	4	45	46	47	48	64	20	51	25	53	\$	55	20	24	28	29

Table 2-6: Poor vs. Good Clinical Outcome: Genes upregulated 5 fold or more in at least 2 tumor samples.

Table 2-6: Poor vs Good Clinical Outcome: Genes upregulated 5 fold or more in at least 2 tumor samples

# of Tumors	, , , , , , , , , , , , , , , , , , ,
Fold(mpm34/ AveGood)	23.80 62.14 0.86 10.17 7.73 6.80 8.85 7.64 6.75 6.98 6.70 5.86 5.81
Fold(mpm33/A veGood)	34.98 0.37 12.26 0.77 1.51 0.22 0.03 0.75 2.47 6.19 6.00 0.66 0.26 0.36
Fold(mpm32/ AveGood)	67.56 62.30 25.33 10.19 7.75 6.82 8.87 7.66 6.76 1.58 1.26 5.87 5.82 5.32
old(mpm31/Av eGood)	2.52 0.24 0.81 0.82 3.69 4.12 0.02 0.95 0.95 1.06 1.37 1.66 0.37
Fold(AvePoor/A Fold(mpm31/Av Fold(mpm32/ Fold(mpm34/ veGood)	29.71 31.26 9.81 5.49 6.17 4.49 4.25 3.95 3.83 3.51 3.19 3.15
AvePoor	42.16 19.98 8.54 1.50 3.29 108.55 237.74 3.63 2.37 16.09 7.78 5.16 2.09 3.83 169.35
AveGood	1.42 0.64 0.87 0.27 0.64 24.18 53.51 0.86 0.56 4.08 2.03 1.47 0.60 1.20 53.79
Tissue Distribution	thymus prostate brain spleen testis bone liver testis bone colon heart testis bone colon heart testis bone
Accession No.	AA436479 AA700876 W93315 N76201 W74079 AA664101 AA67706 N58558 H69148 AA432030 AA432030 AA432030 AA432030 AA478298 W84761
IMAGE Clone Id	753076 452374 357105 284601 346321 855624 460487 248412 212188 782513 782513 362059 454822 415766 740941
Order	- 7 c 4 c 0 k 8 c 0 t t t t t t t

Table 2-7: Good vs. Poor Clinical Outcome. Genes upregulated 3 fold or more in at least 1 tumor sample.

Table 2-7: Good vs Poor Clinical Outcome: Genes upregulated 3 fold or more in at least 1 tumor sample.

		Gen Bank			E old (Ave Good (A	Fold(mpm27/Av	Fold(mpm28/Av	Fold(mpm29/Av	Fold(mpm30/Av	End Assessed A Fold (mpm27/Ay Fold(mpm28/Ay Fold(mpm29/Ay Fold(mpm30/Ay Fold(mpm35/Ay	*
Order	Clone	Accession Number	AveGood	AvePoor	vePoor)	ePoor)	ePoor)	ePoor)	ePoor)	ePoor)	Samples
		0003011		6	10.3	707	6 80	3.51	0.68	7.30	4
-	299679	870C0M	2.55	0.50) o	5.6	3.61	6.33	4.17	6.26	4
~	811162	A44804/1	28.87	20. 20.	17.4	2 5	5000	4.22	2.10	21.15	က
က	471641	AA034945	11.03	4.08	10.17	21.13	7.4	72.0	9.85	7.08	ო
4	281039	N47717	36.67	4.42	8.29	78.77		790	7.05	8.80	က
10	282089	N48259	27.13	3.45	7.87	22.41	L 4.0) ·	9.50	15.90	m
g	155072	R71393	3.32	0.45	4 .	8.50	76.	9.0	9 6	10 17	er)
7	208210	H65286	2.70	0.42	6.48	14.00	98.0	9 9 9 0	0.0 %	2.5	, w
œ	770066	AA430545	26.72	4.62	5.78	12.05	0.67	5.6	9 5	12 02	· 64
· თ	712950	AA282273	38:22	6.94	5.50	8.84	2	0.35	. 4. o	2.35	om
. e	81331	T60111	4.13	0.82	5.03	11.18	1.30	0.80	0.00 0.00	5.40) (°
÷	287778	N59158	3.49	0.71	4.90	14.39	0.35	0.46	3.80	7 7 6) r
: \$	4122R	R58953	0.05	0.0	4.38	9.07	2.53	4.46	2.14	5.0	, c
ž	1220	T99191	9 6	5 6	4.16	12.66	4.17	0.79	0.0	3.18	? (
2 \$	186048	H43317	20.00		3.93	3.28	1.41	1.33	4 .30	9.31	າ ເ
ţ	404003	AATTRATE	5) . 65	3.87	4.86	2.38	4.54	2.55	5.03	? (
<u>.</u> 4	105365	R88904	18.1	20.00	3.62	6.03	1.45	0.88	3.51	6.23	, c
2 ;	2556	M46743		370	3.43	5.38	4.18	4.79	2.28	0.54	? (
>	41.00	AA456075	27.70	ر د در	9 6 6 7 6	5.71	4.22	4.22	1.32	1.35	(n)
<u>o</u>	90900	NAK282	4. A	4 KK	333	7.31	0.87	1.42	3.24	3.77	(N)
2 6	202180	A 6.75.748	 5. 4.		40. F	5.85	1.71	0.34	4.76	3.74	m (
₹ ;	(40020	7020240	<u>.</u>		200	2.50	3.69	3.70	1.59	3.11	က
5 (/848CL	078014	5.0	5 6	7 60	2 2 2	3.21	3.87	0.79	3.97	ო
22	1034473	AA//9/28	76.12	97.75	, c.o.	4 16	1.66	1.35	3.32	3.28	က
23	138444	7/7902/	202	t !	2.73	3.70	1.20	0.99	3.13	3.31	က
₹:	378488	AA777187	17.01	71.7	44.65	6.60	25.69	28.44	1.49	1.02	7
8	767130	AA424050	4.28) () ()	5.5	. t	7.47	44.45	1.65	1.22	7
9 1	30/33/	077CAV	10.81 10.81	9 3 3 6	80 8	<u>+</u>	1.15	33.58	2 .	3.17	7
7 6	302023	ERRA38	1.4.4	, c	5.03	1.46	5.93	15.74	1.30	0.71	7 1
9 6	201735	N90882	- 4	9 6	4.56	9.05	2.75	2.93	0.99	7.09	71
3 5	136805	R35051	32.22	7.39	4.36	2.55	3.53	11.13	1 .94	2.65	N
3 2	362278	AA001219	5.53	4	3.91	2.64	2.34	0.78	6.56	7.23	7 (
; £	756708	AA443903	2.58	0.67	38.5	0.50	4.59	10.09	2.99	1.01	N 6
: E	28502R	N21084	1.22	0.32	3.78	1.00	4.36	11.96	0.59	88.0	V
3 3	1409509	AA868929	2.84	0.75	3.78	0.36	5.86	11.75	0.59	0.31	7 (
. %	365707	AA025434	48	0.53	3.68	0.58	3.36	10.11	1.36	3.00	7 (
3 %	768953	AA424920	3.43	76.0	3.66	0.96	5.41	10.03	0.93	0.97	Ν (
3 5	357528	W94121	1.80	0.49	3.65	5.62	0.55	0.68	2.00	9.41	7
;	>*> ->>		}	;)						
						•					

1.38 9.44 8.17 2.75 0.88 9.90		0,79 0,72 0,72 0,73 1,24 1,24 1,24 1,24 1,24 1,24 1,24 1,24	2.44 2.27 4.89 2.27 2.27 2.33 3.42	7.00 7.103 7.11 7.26 7.26 7.25 8.89
5.55 6.90 7.24 0.99 5.55	0.1 0.0 0.0 0.0 0.0 0.0 0.0 0.0 0.0 0.0	3.20 0.60 1.70 6.81 6.82 0.82 7.4	- 6.9.9.9.9.9.9.9.9.9.9.9.9.9.9.9.9.9.9.9	1.04 3.15 3.69 5.65 1.21 1.21 1.76 6.54 6.54 0.28
2.07 0.73 1.32 2.38 5.00 10.28	2.26 1.26 0.59 0.91 1.80	0.57 0.74 4.46 0.94 3.16 6.22 1.02	0.79 1.14 1.55 7.31 7.85 1.62 0.54	6.64 7.66 7.66 7.66 7.66 7.66 7.66 7.66
2.52 2.52 2.54 2.54 3.54 3.54	2.36 3.72 2.36 2.36 1.26 0.81 5.84 0.77	2.99 2.99 1.13 3.75 3.46	1.98 0.96 1.25 1.25 1.22 1.46 0.37	2.03 3.72 3.72 3.72 3.38 3.38 3.78
9.19 1.63 1.01 3.72	6.00 6.09 6.07 7.62 6.24 8.86 8.86 8.86	6.94 6.94 6.94 6.94 6.94 6.94	4.84 0.63 0.63 0.63 0.63 0.73 0.74 1.54 1.54 1.54 1.54 1.54 1.54 1.54 1.5	3.55 2.51 2.51 2.51 2.51 2.54 2.63 3.24 3.24 0.31 0.00 0.00 0.00 0.00 0.00 0.00 0.00
3.54 3.53 3.54 3.25 3.29	3.28 3.28 3.13 3.14 3.12 3.12 3.13	2. 2. 2. 2. 2. 2. 2. 2. 2. 2. 2. 2. 2. 2	2 : 2 : 2 : 2 : 2 : 2 : 2 : 2 : 2 : 2 :	, , , , , , , , , , , , , , , , , , ,
7.90 8.55 0.02 4.11 4.11 0.79	3.68 2.46 0.25 1.18 1.88 0.98	0.02 0.75 1.36 0.64 36.00 0.75 0.45	17.94 14.62 1.22 1.09 1.00 1.50	2.46 0.94 1.63 0.32 1.57 7.91 0.48 0.57 1.52 1.66
28.75 30.23 0.08 13.79 2.60	12.10 8.05 0.83 3.79 7.05 5.91 3.06	0.06 2.32 4.17 1.82 101.29 2.12 98.80 9.80	2.29 48.94 39.72 3.31 2.93 1.50 3.92 3.92	6.38 2.44 4.21 0.81 19.49 1.87 3.37 1.16 52.42 3.61 3.93
R45192 R77948 AA443967 AA633658 AA412417 R66101	H09076 R00276 R61372 AA628113 N5347 AA775257 T67128 AA432270	H47076 H02837 AA418748 R45517 N55459 N59816 AA196979 AA700680	H92821 AA487902 T56281 AA878120 T69477 AA496988 N53031 AA150500 T69305	N74131 H17121 AA702335 R61883 H72722 N34316 AA634526 AA63431 R54590 H53340 N23139 AA479795
35058 145740 757200 856575 731469	46166 123264 37980 1055707 245277 878596 66599 782335	193087 151477 768020 35329 245990 288983 646037 434952	231574 840590 78353 1418621 66902 823655 246430 491751 82905	298417 51178 447416 42872 232772 277173 795425 743868 40100 202535 267546 740604
885444	3 4 4 4 4 4 4 6 2 2	\$ \$ \$ \$ \$ \$ \$ \$ \$ \$ \$ \$ \$ \$ \$ \$ \$ \$ \$	0 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2	62554555555

Table 2-7: Good vs. Poor Clinical Outcome. Genes upregulated 3 fold or more in at least 1 tumor sample.

3.86	1.24	4.33	404	9 5	\$ \ * 0	3.43	0.90	1.87	4.40	3.90	1.32	5.80	4.05	4.08	4.96	3.89	4 53		9 6	0.92	3.62	3.44	4.38	3.80	1.03	3.61	35.6	3.05		2.5		9.00	, c	- 67	3.10	2.03	4.12	47.4	4.0	3.03	10.0	3.76	3.15	1.1	0.61	0.49	0.48		
2.42	3.44	900	9 6	9.00	0.18	4.10	0.74	4.69	0.24	4.70	3.63	0.51	5.78	9 6	20.0	9 4	3 -	- G	9.00	 	1.67	2.03	3,46	1.07	106	77	1 8		S. 50	5.4.	4. c	91.9	3.09	0.83	3.41	3.03	0.54	0.58	3.49	0.41	0.75	3.65	3.37	3.19	0.56	0.63	4.04		
0.76	· .	7	20.0		0.08	0.30	3.31	0.42	0.11	16.0	2.5	9.45	? ?	2 4	6 C	0.40	20.	0.97	0.49	3.40	3.00	0.93	1.22	75	3 0	2.02	9.	7	0.41	1.13	0.52	0.98	1.23	0.68	0.93	0.78	0.37	0.38	0.85	1.87	4.05	0.52	0.21	3.72	3.89	3.81	3.25		
4 70	7.	1.5/	o 4	1.55	2.10	0.65	1.49	0.83	5.59	5	5.5	0.02	9 7	rc.0	7.00	4.41	0.97	3.49	0.49	3.05	1.35	0.50	0.67	9 0	20.0	78.1	3.63	2.55	3.29	3.70	0.58	1.69	1.05	1.99	0.93	0.75	3.84	3.02	0.90	3.19	3.33	99.0	7.	0.56	3.19	3.18	0.53		
ç	0.50	9.94 49.	3.32	3.12	4.65	2.81	4.73	722	0.57	9 9	0 0 0 0 0	90.5 6	3.40	0.17	0.32	1.03	3,28	4.01	0.34	1,59	0.89	3.61	97.0	0.0	96.0 00.0	3.33	0.84	3.08	1.09	0.56	0.27	0.76	1.43	3.13	1.39	1.40	0.92	1.25	0.74	0.91	0.55	0.42	0.69	0.26	0.57	0.36	000	; 6	っ
	2.36	2.32	2.31	2.30	2.27	2.26	2.23	3 6	2.52	77.7	2.22	2.20	2.19	2.19	2.18	2.17	2.17	2.14	2.13	2.11	24.0	- Ç	2.70	2.70	2.09	2.07	2.06	2.06	2.04	2.01	1.99	1.99	1.98	1.97	1.96	1.96	1.96	1.89	1.89	1.89	1.86	1.80	1.77	177	1 76	2 5	8	2	
•	1.33	0.33	<u>5</u>	0.37	6.85	1 26	0.50		4) (4	14.0/	5.54	40 0	17.15	15.57	13.39	2.98	0.24	1.57	10.50	1 71		3 5	44.4)	0.86	0.30	0.83	10.94	9.27	3.81	0.01	25.44	2.07	1.1	1.50	0.85	15.	2.19	8.31	0.76	15.04	13.84	3.24	21.21	27.34	7 24	700	; ;	0.0¢	
	3,15	0.77	2.41	0.84	15.54	300	£.03	<u>.</u>	8/.4 8/.5	28.11	12.27	3.07	37.52	34.05	29.18	6.47	0.52	3.35	22.38	3.60	9 6	9 (c)	90.74	18.	0.62	1.73	22.57	19.09	7.78	0.02	50.71	4.12	2.20	2.96	1.67	2.97	4.29	15.71	1 44	28.35	25.74	5.85	37.58	48.33	57.3	2.72	B C	3	
	AA021586	N67039	AA447522	H08862	T67053	NE2464	A A 45.6300	000000	AA456321	AAGGSBBI	AA988345	R76247	H16803	H77766	N57872	AA884926	H50086	AA434487	R16596	W49563	200011	A A G 20006	AAU/0220	AA779457	H15677	AA609566	AA111969	AA668897	AA683520	AA459292	N80129	AA035558	R36989	AA459197	AA040424	N64706	N92646	70057	AA457528	W70234	N30680	AA489666	AA630006	R27004	4448000	DA6000	AA80004A	71000011	
	364022	295985	782575	AKRKO	10001	00000	284505	101010	813179	855745	1605539	144747	50772	214162	247117	1468260	179083	770868	120KR	201046	254013	/80L8Z	530814	1032405	49303	1031701	530185	853998	378813	810899	297392	471682	27098	810873	485854	293292	289337	80048	838500	343087	258300	824340	884606	432702	707740	10/4th	40000	5111501	
	æ	젏	8	ä	3 8	5 6	8 8	8	8	5	85	8	¥	82	8	26	88	8	\$ \$	3 5	5 5	202	ន	Ž	105	8	107	108	60	5	=======================================	÷	. <u> </u>	7	- -	£ 5	÷	. ¥	5 5	5	į	÷	<u> </u>	3 5	5 5	07 9	9 5	12/	

Table 2-7: Good vs. Poor Clinical Outcome. Genes upregulated 3 fold or more in at least 1 tumor sample.

2.25 2.00 2.00 2.00 2.00 2.00 2.00 2.00	1.55 9.72 1.25 1.25 1.16 1.00 1.24 1.37 2.55 2.55
1.75 1.77 1.00 1.00 1.52 1.30 1.78 1.78 1.78 1.78 1.78 1.78 1.78 1.78	2.72 0.32 0.32 1.53 1.53 1.53 1.53 1.53 1.53 1.53 1.53
112.76 0.85 29.22 29.22 2.0.07 2.10 0.99 15.25 1.03 10.97 12.03 0.30 0.30 0.30	6.33 0.63 0.63 0.63 0.63 0.63 0.75 1.38 1.38 1.39 0.72 0.82 1.38 0.82 1.33 1.33 1.33 1.33 1.33 1.33 1.33 1.3
29.27 0.57 0.57 1.05 1.05 1.05 1.05 1.72 1.72 2.24 1.73 2.24 1.53 2.78 2.24 1.53 2.08 1.53	1.67 0.91 0.94 1.37 1.65 1.65 1.65 1.66 0.82 0.60 0.60 0.45 1.03 1.03 1.03
0.70 0.79 1.41 22.37 0.45 0.65 1.25 1.57 1.57 1.57 1.50 1.50 1.50 1.50 1.50 1.50 1.50 1.50	2.06 0.85 11.85 9.16 2.20 7.19 7.19 7.19 8.73 8.73 8.73 8.73 8.73 6.15
66 66 66 66 66 66 66 66 66 66 66 66 66	2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2
0.85 0.69 0.1.1.23 0.50 0.50 0.50 0.50 0.50 0.53 0.53 0.5	7.56 7.40 7.40 7.56 7.53 7.65 7.65 7.65 7.65 7.65 7.65 7.65 7.65
4.56 8.01 2.77 4.56 6.75 4.47 1.51 1.93 13.18 1.95 11.01 2.103 16.43 17.91 17.91	45.4 45.4 45.4 45.4 45.4 45.4 45.4 45.4
N78927 N48325 AA702640 N90806 AA677574 AA677574 AA677574 AA677574 AA677574 AA677574 AA677574 AA677574 AA677574 AA677574 AA67757 AA467608 AA680136 AA680136 AA680186 AA680186 AA680186 AA680186	H52001 W81196 R00284 AA452113 AA398348 H29897 H20046 AA018338 H45810 H11433 AA504842 AA504824 AA504824 AA504824 AA5013094 W86202 N29545 AA620757 N68719 H80853 AA680070 T99817 R44664
300051 279690 384015 303139 455271 844703 841507 427978 275738 258063 141281 795265 133130 611581 812105 430465 1637302	197525 347429 123265 786525 726835 190325 172786 362419 188151 47652 825736 825770 258242 309893 360213 416374 416374 416374 430368 123354 34129
821 821 822 823 824 825 827 827 827 827 827 827 827 827 827 827	149 150 151 152 153 154 155 155 155 156 167 168 168 168 175 175 175 175 175 175 175 175 175 175

Table 2-7: Good vs. Poor Clinical Outcome. Genes upregulated 3 fold or more in at least 1 tumor sample.

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	1.31		9. 48.	2.39	0.96	1.95	1.55	3.34		7.07	0.00	2.09	1.42	0.68	7.52	1.41	1.48	109	53	- c	7.10	0	1.63	0.91	1.23	2.71	1.52	0.85	5.74	3.00	1.25	55		2 57	; 6	20. PC) c		2.6	9 5	2 5	5.6	\$ 1	5.7	71.1	1.62	1.26	1.32		
	0.60	2,93	0.81	3.69	0.87	1.37		3 7	* (*	5.73 5.73	0.62	3.29	2.65	0.58	0.68	0.50	44.	5	- c	9 1	7.7	2.96	1.45	0.73	1.20	0.99	2.58	0.86	171	2.47	800	20.0	2.4.	 	¥7:7	, c	77.7	4.40	0.00	2.13	2.03 20.03	0.70	46.0	3.43	2.	2.30	0.75	2.72		
÷	6.37	1.21	5.79	1.60	60	5 6	5 5) (i)	2.59	1.31	2.07	2.31	5.23	2.79	1.29	17	. 2	9 6	00.	2.14	1.47	2.97	5.01	1.21	5.81	0.60	4.41	88.8	8000	9 6	9.00	- C	0.92	2.37	77.	90.0 10.0	5.5 5.5 5.5 5.5 5.5 5.5 5.5 5.5 5.5 5.5	S	07.1	2.07	1.50	1.15	7.24	2.25	1.50	3.09	, 2	1.89		
	1.73	2.07	2.76	1 78	- 6	0.00	0.87	1.30	2.05	2.57	8.64 4.64	1.79	1.49	2.27	0 74	2.44	; ;	3 6	2.83	0.91	3.68	1.57	2.63	0.66	2.75	1.47	5. 4.	20:- 0	6.95 4.36	07.	9.60	2.78	0.77	3.50		0.87	2.5	0.92	2.38	2.45	4.49	0.96	1.0	1.25	1.78	2.64	2.95	1.56		
	2.49	5.32	2.25	3 6	7.04	J. 58	6.85	1.73	2.69	5.71	0.42	2.81	7 78	. v	9 6	2.00	0.0	2.70	6.14	5.00	2.98	3.34	1.30	8 49	1 00	9 6	0.40	3 5	0.40 0.40	2.79	1,31	1.87	6.37	1.24	3.84	7.58	1.80	3.82	4.34	1.35	2.91	7.79	1.54	2.87	60.9	1.86	5.46	3.95	· ·	>
	2.50	2 50	3,40	D (*)	2.48	2.48	2.48	2.48	2.47	2.47	2.47	2.46	27.6	2,43	7.43 7.43	2.40	2.44	2.44	2.43	2.42	2.41	2.40	2 40	2 70	9 6	7.40	8.39	25.33	2.38	2.38	2.38	2.38	2.37	2.37	2.36	2.36	2.36	2.35	2.35	2.34	2.32	2.32	2.31	2.31	2.31	2.30	2.29	2.29		
	0.19		9.6	64.7	1.86	0.98	2.05	0.25	1.02	0.02	1 70	20.4	77.	1.48	51.01	3.12	0.05	2.13	0.82	0.31	08:0	1 22	1 2		7.7	ان 10.0	0.87	3.48	0.61	4.19	2.90	1.89	1.57	9 8	12.22	0.42	2.54	0.67	0.62	90.0	9.0	0.29	0.59	2.31	1.68	8.14	35	2.22	*	
	0.47		(9.10	4.61	2.42	5.08	0.63	2.53	90	4 20	7.50	- 6	3.62	24.84	7.63	0.12	5.18	1.98	0.74	2.17	203	3 6	3 5	0.0 1	1.45	2.08	8.32	. 4	9.95	6.83	4.49	3.73	90.0	28.85	6 .0	5.99	1.57	1.46	0.19	0.10	0.68	1.37	5.33	3.83	18.76	02.0	2 c	Š	
	AA455654	070270	**************************************	かかつくつせる	AA953357	AA609218	H09959	AA486561	AA287122	AA262080	AA875913	AA48443	24040	K08980	KSSST	N5/002	AA620783	AA455087	AA444083	AA039851	AA707086	AA431887	A A D 3 5 4 5 0	U30637	1/85/2	K48/31	N72115	T64312	AA029578	AA136618	R39325	AA190789	AA425947	AA666348	AA045698	R63085	AA972352	H15099	N50853	W74653	N33264	R38891	AA872006	AA405543	R27193	H69691	NO1285	WARRES	***	
	844004	10010	910806	757337	1573251	1031478	46367	840967	701819	685801	4402202	1432202	810328	128738	26736	277596	1049321	812294	756657	375827	451587	773630	113039	67/1/40	739611	38554	291057	83920	366859	490946	24176	627351	760299	858849	509731	137984	1584628	49354	280957	346712	273501	25052	1475648	772416	133341	212784	206043	416833	410000	
	Ş	2 ;	174	175	176	#	178	179	£ 5	5 5	5 6	70 5	2	≅ ;	185	186	187	188	189	8	Ş	: :	7 5	2	\$ 1	195	1 8 1	197	198	199	50	201	202	203	8	205	206	207	208	208	210	211	25	<u> </u>	2 2	2 4 4	2 6	6 4 6 7 4 6	/17	

Table 2-7: Good vs. Poor Clinical Outcome: Genes upregulated 3 fold or more in at least 1 tumor sample.

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1.46 6.91	1.85	. 790	121		3 5	 	07:	5.T	05.7	0.95	1.35	3	4.78	1.92	<u>.</u> 26:	4.63	1.95	3.44	4 20	2 7		1.32	6.18	0.91	5.06	2.32	2.13	0.78	2.22	27.0	÷ 60	16.0 78.0		1.94 200	7.00	2 6		2 2	2.24	1.5	2.08	1.02	2.31	9.0	0.63	1.14	· •		
2.16	90	5 6 6	5.65	3.5	.83	1.73	<u>.</u> 8.	1.23	1.52	1.14	2.63	2.03	44.	1.98	1.67	200	4 55		2 6	2.07	1.89	0.67	1.66	0.63	1.38	20.5	234	2.4	<u> </u>	 	2.20	5.00 00.00	30.5	0.93	0.75	5.7	3.13	2.02	1.32	1.07	1.75	0.93	1.74	0.67	4 33	10.0	6.65		
0.42	3 .	4.17	0.90	. 4	3.50	3.19	5.54	0.76	5.46	5.42	4.12	88	25.	- T	9 6	2.7	2.11	\S.4.	0.79	2.87	4.67	1.83	760	6.28		6.6 7.6	 	41.1	5.62	1.78	0.93	1.05	3.25	4.58	1.03	 \$	2.39	2.76	1.64	2.45	6.00	1 60	3.22	1 d	0.0	4 .	1.49		
6.23	,	2.73	2.76	1.95	1.95	1.95	1.85	0.71	50	67:1 VV C	7.5.4	è F	27.4	7.50	1.59	0.89	2.11	0.72	1.97	1.19	2 13	- 4B	5 · ·	17:1	2.3	0.60	1.51	<u>.</u> Ş	1.83	1.74	2.25	1.64	2.63	1.61	1.02	1.78	1.42	1.12	2.37	1.76	96	3 6	5.5	.c.	1.43	0.71	2.02		
1.11	0.84	0.61	5.34	5 62	2 4 7	. t t	9 6	1 0	C7: /	40.	81.1	1.33	1.45	1.06	96.0	3.76	1.15	1.90	2.71	, r	3 6	5 r	20.0	0.82	0.67	1.76	1.13	4.21	1.05	2.03	4.70	2.48	98.0	1 70	7.11	2.42	1 25	1 54	, c		2.70	74.	5.74	1.78	1.00	7.07	5.29	9	
2.27	2.27	300	3 6	2.43	27.7	2.25	2.25	2.24	2.23	2.23	2.22	2.22	2.22	2.21	2.21	2.20	2 19	5 40	. c	6.19	2.18	2.18	2.18	2.17	2.17	2.17	2.17	2.16	, t		7. c	, c	2.13 4.13	2.13	2.13	7 C	2 5	2.5	51.3	2.13	2.13	2.13	2.13	2.12	2.12	2 12		- •	
0 0	5 6	9.90	07.1	0.92	0. 48.	0.15	1.05	6.24	0.70	0.42	0.20	2.64	12.29	3.83	1 75	2.0	. 6	3 6	300	9. .	0.65	0.42	3.12	0.29	0.43	0.24		9 6	2.30	4.0	0.0 0.0	0.92	40.0	1.67	0.63	0.67	95.4 90.4	4.40 0 :	0.57	29.69	0.07	0.97	1.63	2.88	2		† ¢	3	
4	3 6	2.03	3.97	2.07	1.89	0.35	2.35	13.95	1.56	0.95	4	88.	27.25	() () () () () () () () () ()) d	9 6	70.1	4.0	0.19	2.86	.	0.92	6.81	90	. 6	3 5	70.0	5 6 6	6.27	8.5	0.13	1.98	1.16	3.59	1.35	- 4	10.58	9.38	1.21	63.20	0.14	2.07	3.46	11	- ç	2.0	2.41	7. 1.	
900	610010	NZSB14	AA044906	H40921	AA703652	N66572	AA481045	H54417	AA460422	AA676265	H10342	72067	AA703396	D00884	A449962	70000	AAU34133	AA663092	H08397	AA487560	AA055221	AA005350	H10036	W02426	A A 46/12	A 4 4 3 2 2 5 3	74456635	K381/9	/9LGCN	AA457138	H19324	H72093	AA608575	AA137078	H02039	AA626275	AA708438	AA455151	H15408	AA156054	H12081	AA706738	AA487468	A A R 20003	AA023903	AAU AU AU	AA056013	T60063	
	430336	271050	488706	177074	450382	278808	814662	203003	796480	424663	45155	0000	82800	4000 d	123971	566440	380437	852577	45501	841664	377368	428377	16630	206170	271087	795755	782306	23869	245547	810459	50922	213604	950710	491196	151067	745072	506011	809881	49227	590338	47597	422000K	044634	170140	884662	383199	377692	81320	
	218	219	220	22	222	3	3 2	, , , , , , , , , , , , , , , , , , ,	3 2	9 6	ž	97	87.5	8	ន	232	233	5 3	235	236	23.7	, a	200	2 2 3	240	241	242	243	2	245	246	247	248	249	250	251	252	253	254	255	35	2 6	2 2	907	259	2 8	5 61	262	

Table 2-7: Good vs. Poor Clinical Outcome: Cenes upregulated 3 fold or more in at least 1 tumor sample.

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0.82	0.26	1.28	1.40	162		7.0	3 6	3.82	1.07	1.08	4. 20.	3.33	1.12	<u>.</u> 2	0.65	2.59	1.15	1.68	1.17	1.11	2.13	2 6	2.0		 	3.81	04.1 0 i	0.95	3,19	90.0	2.65	1.00	2.05	0.95	0.83	21.1	24.0	2.81	7.83	2.19	1.81	1.46	1.49	3.27	0.85	1.86		
0.79	0.61	4.75	9.0	5 6	67.7	2.78	8.00 0.00	76.0	0.96	4.45	1.66	2.07	0.90	66.0	4.59	1.33	0.75	99'0	158	1 78	2. c	9 6	0.60	1.24	1.12	49.	5.35	0.42	2.47	1.40	1.57	1.19	1.05	1.43	0.83	3.90	5.54	1.51	0.60	2 .	7. \$	1.36	0.82	1.70	0.77	4.12		
7.10	60.0	8 6	9.6	0.00	70.4	1.13	0.31	5.69	4.22	0.98	1.51	0.80	1.61	2.17	0.55	0.71	2	0.67	2 S	20. 4	<u>:</u>	\$ i	2.32	5.31	3.05	0.31	1.03	1.98	1.40	2.18	1.40	4.79	1.13	1.18	0.68	1.17	1.27	0.86	1.29	0.86	2.08	1.67	0.48	0.49	0.95	1.76		
96'0	9 0	2 9	24.7	1.42	1.12	2.24	1.48	1.33	2.98	1.40	1.50	141	1.40	1.29	2.20	53	2.78	80.0	3 6	9 6	70.4	4.24	1.91	1.20	1.80	1.58	0.65	4.68	2.37	4.	1.10	2.47	3.34	1.83	1.03	2.36	1.24	1.07	1.43	2.10	1.03	1.65	1.88	1.82	2 .	0.75		
0.88	9.0	9	2.07	6.43	1.07	0.00	0.00	1.64	1,22	2.45	1.15	2.74	. v	A 25.5	200	17:7	7.53	4.04 4.04	0.4		1.39	1.18	4.21	0.98	2.47	2.88	1,78	2.54	0.73	4.	3.43	0.68	2.57	4.75	6.75	1.57	1.63	3.84	4.94	3.59	3.32	3.95	5.41	2.79	5.85	1.57	7	
2 44	- 4	2.10	2.10	2.10	2.09	2.09	2.09	5.09	2.09	2.07	207	2 6	. c	2.0	2.07	7 .00	9 50	9 9	8.5	5.06	2.06	2.05	2.05	2.05	2.05	2.04	2.04	2.04	2.03	2.03	2.03	2.03	2.03	2.03	2.02	2.02	2.02	2.02	2.02	2.02	2.02	2.02	2.02	2.01	2.01	2.01		
73	200	1.29	0.96	0.86	1.76	0.0	0.01	0.40	0.45	1 10	200	7	5.0	8 3		5.4.3 14.0	, to	0.24	1.73	5.34	9.0	2.52	14.49	0.26	0.53	7.71	1.40	0.60	64.19	0.65	0.28	0.38	1.82	0.43	0.97	1.91	0.40	0.62	0.10	3.42	108	0.03	5.26	1.73	1.07	2.74	.	
č	17:1	2.72	2.02	1.80	3.68	0.09	0.02	0.83	98	2.46	9 6	9 9	6.0 0.0	25.1	0.80	5.02	S. S. S.	0.49	3.57	10.99	0.08	5.18	29.68	0.53	1,08	15.75	2.86	122	130.51	1.32	0.57	0.77	3.68	0.87	1.97	3.86	0.81	1.24	0.19	6.91	2.14	0.07	10.60	3.49	2.16	5.46	;	
169567	10000N	AA935560	W96134	T56726	AA485352	AA292429	W72749	AA757717	AA699573	W96155	128740	A 4 500 10	A 1303 10	NO2403	74404L	6187/1	CHACOL	10/153	AA424045	AA775616	AA443602	AA700054	AA777670	AA677050	AA015959	AA996131	H26426	T54164	T64469	R51535	T99852	AA461311	AA463635	AA057156	N48050	H93381	H04382	T61050	AA071503	AA521247	H08424	AA449762	H63416	T64223	H93335	A1004169		
	71877	1556526	358531	67092	810998	714498	346119	396098	433481	368534	10000	49000	Renene	288881																			811827	489079	281659	241821	149373	76988	366039	827152	45823	785816	200025	80221	242061	1631807	; }	
Ç	3	7	265	266	267	268	269	270	274		717	273	4/2	2/2	5/2	27.7	278	279	280	281	282	283	284	285	286	287	288	289	8 8	29.	200	293	26.	282	296	297	298	299	300	301	305	303	Ş	5 5) Ş	3 %	;	

Table 2-7: Good vs. Poor Clinical Outdome: Genes upregulated 3 fold or more in at least 1 tumor sample.

-	-	-	٠ •		_	-	-	-	· •		- ,	- •	-	-	-	-	-	•	- •	- ,	- '	-	-	-	-	• •	- ,	- ,	-	-	•	-	-	_	•	-	_	-	Ψ-	-	•	- *	- •	- •	-	•	_	_	•			
440	2. c	5 6	2.0	0.80	1.1	1,31	4.66	28.0	5 -	t :	40.0	0.97	0.95	1.01	1.14	4 84	5 5	2.5	20.	0.93	4.11	4.99	1.41	230	6.5	9.6	2.14	0.68	0.60	0.88	2.84	2.81	1.37	75.4	, 60 7	20.5	 	5 5	. +	6. 6.	S .		3,35	1.63	2.46	2.18	1.50	25.6	יייי אמי	3		
į	- 6 2. 9	D.87	1.42	0.87	1.56	52	5 5	20.1	1.41	4.19	0.75	0.64	1.48	4 53	3.5	70.1	X	0.53	0.61	0.73	2.22	1.50		6.63	0.81 1	0.70	2.02	0.80	0.45	4	÷ ;	4 60	67.	2.5	S	06.1	1.70	0.80	7.52	0.93	2.79	0.71	2.82	0.93	0.14	1 22	- C	. c	[A]	0.82		
	3.33	0.62	5.26	561		0.00	2.15	2.39	4.42	1.08	0.25	5 14	· 6	3.00	3.80	2.54	1.29	6.33	0.70	66 U	0.44	F •	04.	3.71	2.17	0.50	3.33	6.67	9 6	2 6	3.4 3.4	0.85	0.94	1.70	0.41	3.88	0.39	0.85	3.47	2.39	1.27	0.93	1.58	78	- 6	900	.03 0.03	2.55	1.62	0.59		
	1.27	2.08	18)	1.06	3.32	1.35	1.87	+		70.7	4	2.31	1.96	1.42	1.82	1.17	÷	- 0	S	84.	0.85	2.60	0.88	2.73	2 2	† 6	5.0°	4.60	0.96	0.95	1.53	1.76	1.45	1.59	1.96	1.12	1.49	3.44	0.57	2.45	2 6	5.95	1.01	2.89	1.53	3.05	1.15	0.52		
	2.43	; c	20.0	0.81	0.99	5.19	1.73	, e	2 7 7	÷ c	2.04	2.98	1.75	1.38	1.53	3.29	3.41	- c	5.6	6.42	5.23	1.57	1.03	1 44	99.6	9.6	1.8.1	1.26	0.90	1.71	1.80	0.98	3.17	6	2.17	98.0	2.64	55	2.24		99.	5.00 9.00	50.0	0.98	3.44	4.07	5.	1.76	2.42	88	ζα	כ
	Č	7.01	2.07	2.01	2.01	000		0.00	2.00	2.00	1.99	1.99	1.99	00	5 6	9. 0	68. 68.	28.	1.98	1.97	1.97	1.97	70,	6.	0 8	1.96	1.96	1.96	1.96	1.95	4 0 7	5 4		6.	<u> </u>	<u>.</u>		<u>.</u>	. 3		5.	1.93	1.93	1.93	1.93	1.93	ę	8 6	26. - 1	3.5	? ~	
	1	4.73	2.33	0.92	0 63	1 6		0.46	11.14	3.86	9.	10.17	3.5	9	0,00	77.	3.01	5.09	0.48	3.29	1.52	23.0	7.00	3.2	0.28	6.16	2.46	1.06	1 24		5 5	2 !	2.4/	7.09	0.43	2.22	. .	5.12	0.33	1.	0.11	2.17	0.90	4.52	0.08	47 R4	ţ 8	9. ·	4.49	9.0	1.58	
		9.52	4.69	1 85	3	47.	ž.	0.92	22.32	7.73	60.0	20.04	20.54	0.70	0.72	2.42	5.97	4.15	0.95	6.49	000	7.93	4.5/	6.31	0.54	12.08	4.83	000	3 5	7.70	0.02	2.31	4.82	13.82	0.85	4.32	2.02	9.8 26.0	0. 26	2.19	0.22	4.19	1.74	8 72		2 3	\$ 5	0.12	8.66 9.66	1.24	3.05	
		AA708440	H38650	W15263	00000	H84309	N38860	AA633866	AA630771	H84871	H15417	L777307	167/14	AA404293	AA464935	AA017133	AA421270	170892	W86987	A A 453015	144067	1000 L	N896/6	AA486435	R42520	H42679	R96827	0 8 427740	A A O 4 O 4 C	AA01040	AA443648	R34331	AA453997	AA962236	H23137	AA009609	AA128407	AA055835	AA707872	AA004835	T64216	W86653	R24451	AA703387	A A 425825	77173073	H18423	AA777700	AA121836	R08266	N91914	
		506016	100732	1807.32	32233/	249526	279905	858188	856420	249603	70007	55584	233464	758366	838839	362628	731031	83999	44EKKE	10000	10007	163841	307053	842918	29954	183337	10000	1007	770837	362424	771258	136474	795262	1602120	52303	365515	526587	377461	413150	429148	80226	416833	122054	10000	95005	548693	50877	449257	566106	127193	306829	
		308	3 6	9 C	310	311	312	343	314		2 6	316	317	318	319	320	32	2	1 6	253	324	325	326	327	328	200	R70	250	331	332	333	334	335	338	337	338	339	8	3	342	72	3		Ç,	84.0	7	3 5	378	35	351	352	

Table 2-7: Good vs. Poor Clinical Outcome. Genes upregulated 3 fold or more in at least 1 tumor sample.

2.28 2.29 2.20 2.20 2.20 2.20 2.20 2.30 2.30 2.30	2.52 0.93 1.14 2.41 1.36 1.36 1.16
2.1.1.2.2.4.2.4.2.4.2.4.2.4.2.4.2.4.2.4.	3.50 1.12 2.02 3.32 0.97 0.97
20.65 20	0.31 0.88 0.72 0.99 1.54 1.58
0.11 0.82 0.82 0.94 0.94 0.95 0.95 0.95 0.95 0.95 0.95 0.95 0.95	0.63 1.28 0.89 1.07 3.75 0.65 1.31
0.94 1.90	2.37 2.18 3.17 1.16 1.70 1.70 9
2. 1. 1. 1. 1. 1. 1. 1. 1. 1. 1. 1. 1. 1.	2. 1. 1. 1. 1. 1. 1. 1. 1. 1. 1. 1. 1. 1.
1.99 1.45 1.45 1.45 1.55	25.55 1.25 1.81 2.63 0.01 0.39 0.57
3.83 2.74 2.75 7.48 9.09 9.09 9.09 11.69 12.38 12.38 12.38 12.38 12.38 12.38 13.08 14.11 14.11 15.33 16.69 16	22.84 1.88 3.38 4.89 0.02 0.73 1.72
H41489 AA883735 AA426113 AA426113 AA426113 AA426113 AA460827 AA666418 AA460827 AA666418 AA666418 AA66627 AA666418 AA66627 AA66627 AA66627 AA66627 AA666418 AA68576 N50912 N91003 AA68526 N73301 AA676434 AA776434	AA775447 AA775447 AA931725 AA626315 H52361 R43535 AA707840 R26859
192569 1466788 1573305 757285 687667 878281 589362 796288 859654 488301 61412 45877 841615 281072 306066 510576 431286 214809 269425 592243 1555451 292068 825356 45376 727764 39933 290235 11832 396045 701272 1471841	1471841 378182 307660 1572196 745570 202168 32578 412949 133358
353 354 355 355 357 357 357 377 377 377 377 377	330 330 337 337 337 337

Table 2-7: Good vs. Poor Clinical Outcome: Genes upregulated B fold or more in at least 1 tumor sample.

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1.25	3.19	0.97	4.40	0.95	0.86	1.48	185	- C	6.53	3.	1.1/	1.31	2.44	1.73	0.63	0.84	0.98	4.	1.38	000	1 10	- 4	10.1	1,3/	0.95	2.35	<u>.</u> 2	0.81	1.02		0.99	0.97	2.01	1.20	1.24	3. 3.	9:00	5.40	1.39	1.09	3.55	1.01	0.98	1.20	1.73	1.56		
1.16	2.54	1.45	16.0	1 16	52		9.60	8.5	1.26	0.80	0.89	1.22	1.72	4.47	0.57	1.67	198	3.16	; ,	3 6	9.6	9 9 9	1.7.	1.68	0.88	1.21	1.38	0.58	1.01	3.04	1.73	3.31	1.21	0.56	2.21	0.73	1.18	1.28	1.79	1.06	1.05	1.20	0.80	1.73	1.17	1.75		
5.01	0.48	7 9	4.1.	2 4	4. 4. 4. 4. 4. 4. 4. 4. 4. 4. 4. 4. 4. 4	4	0.99	1.00	4.60	3.77	1.02	0.86	3.18	1.42	89	5 6	- c	 	71.7	7.07	4.94	4.13	3.33	3.09	4.14	99.0	3.26	4.97	3.19	2.29	4.08	1.99	0:30	3.28	3.14	1.11	3.59	0.79	1.37	4.06	1.95	3.63	2.16	2 7.3	9 6	9.40	2	
٠ د	3 5	71.1	1.17	5.	1.25	0.84	1.50	3.33	0.92	1.72	1.17	2.24	10.4	7: +		P (C	77.	1.26	1.30	1.96	2.28	1.40	1.22	105	48	5. 5.	1 76	5	2.55	76	1.26	88.0	0.24	0.98	1.43	1.03	1,35	0.93	12.0	. 4	2 7	<u> </u>	 85.	2 6	1.82	2.3	<u>†</u>	
7	5.73	1.91	1.48	1.40	1.33	1.02	3.24	2.16	1.85	188	20.7	 	5.50 5.00	0.62	0.43	2.49	3.52	1.86	1.12	3.69	1.10	1.33	5	- •	5.5	77:-	0°.0	. c	0.90	t	† 60 - 0	0.90 78	0.1 ac a	03.0	66.7	3.20	23.0	2.50	6.0 84.0	5.00	7:01	9 . 9 !	7.5	3.23	1.35	3.48	0.58	10
1	1.85	1.85	1.85	1.85	1.85	- 48.	48	8 4	5 %	6.0	ę ;	\$ 3	2	1.83	1.83	1.83	1.83	1.83	1.83	1.82	4 82		70.	7.87	7.87	1.82	1.82	1.82	£	1.81	1.81	£0.5	S .	8. 8	 8 6	9 6	9 8	 	 8	8. 1	1.79	1.79	1.79	1.79	1.79	1.79	1.78	
	1.27	15.36	0.87	0.12	1.27	990	4 47	F C	7.07	0.80	0.56	0.25	0.46	0.78	0.11	1.16	0.71	1.09	68	0.34	5 6	3 9	D (1)	0.80	1.66	37.83	2.47	0.09	0.87	7 .8	9.31	0.26	1.35	2.56	0.95	15.33	0.80 0.80	1.39	0.39	5.49	0.51	0.45	0.63	14.15	18.60	8.45	10.58	
	2.36	28.41	1.61	120	2.35	1 10		7.7	4.04 4.04	1.47	1.03	0.47	0.85	.43 643	0.21	2.12	1.30	2.00	3.46	9 4 6	9 6	0.0	0.80	1.45	3.01	68.74	4.49	0.17	1.58	13.83	16.82	0.46	2.43	4.62	1.72	27.62	. .	2.51	0.70	9.86	0.92	0.80	1.13	25.27	33.23	15.09	18.88	
	AA704587	AA886870	H22563	AA490987	H19826	A A 699714	A 463400	A4405100	W32408	R80217	AA505045	H17696	AA411387	AA441933	AA868515	AA609049	H20547	AA099582	A A O 1 G A R 2	140761		K48320	AA707935	AA098896	N47604	AA775863	AA872454	AA199881	H99075	AA455042	AA975209	AA458491	AA180204	AA775223	AA521394	AA976544	R97226	R48833	AA682293	AA287404	AA969184	H09747	N32904	AA609474	N35067	H23211	T78909	
	450598	1403252	54016 54016	01010	477547	11.6211	43328/	796876	321359	147050	825648	50043	754601	774078	1408710	1031278	E4420	489706	20000	303000	801/4	153541	392399	489553	277848	878544	1476019	645259	262060	812244	1555427	809620	611412	868838	826072	1558642	200395	153617	461727	701112	1580342	46353	259884	1031598	271662	51773	108864	
	398		800	} {	2 §	Ž (50 3	\$	405	4	404	804	409	410	¥	÷ ;		2 7	* :	415 5	416	417	418	419	420	421	422	423	424	425	426	427	428	429	430	£3.	432	433	434	435	436	437	438	439	94	177	442	!

Table 2-7: Good vs. Poor Clinical Outcome. Genes upregulated 3 fold or more in at least 1 tumor sample.

Table 2-7: Good vs. Poor Cilnibal Outcome. Genes upregulated 3 fold or more in at least 1 tumor sample.

1.19 3.23 3.05 4.02 0.84 0.97 0.98 0.98 0.66 0.66 0.75 0.75	2.5.2 2.2.2 2.2.2 2.5.3 2.5.3 3.68 3.68 3.68 3.73 3.73 3.73 3.73 3.73 3.73 3.73 3.7
1.30 1.30 1.26 1.26 1.26 1.26 1.40 1.51 1.51 1.55 1.33	1.36 1.25 1.09 1.70 1.57 1.37 1.30 1.31 1.31
3.64 0.84 0.93 1.74 0.90 1.12 1.13 1.15 1.15 1.26 0.50 3.79 3.79 3.05 0.50 0.50 0.50 0.50	24.1 20.1 30.1 30.1 30.1 30.2 40.0
1.27 1.00 1.54 0.78 0.93 0.82 1.12 1.38 0.38 0.58 0.58 0.58 1.52 0.58 1.33 0.58 1.33 1.33 1.33 1.33 1.33 1.33 1.33 1.3	0.71 1.20 1.12 1.13 1.14 0.85 0.73 0.69 0.69 0.73 0.73 0.73 0.73
3.95 1.06 0.66 0.66 1.67 1.27 1.27 3.09 3.09 1.19 1.20 1.78 1.20 1.78 3.03 3.63 4.03 3.03 3.03 3.03	3.73 2.63 3.32 3.32 3.32 3.32 3.13 3.15 3.26 3.26 3.17 3.15 3.15 3.15 3.15
	1.70 1.70 1.70 1.69 1.69 1.68 1.68 1.67 1.67 1.67 1.67 1.67 1.67
2.34 2.35 3.30 3.30 3.30 3.30 3.30 3.30 3.30 3	1.89 1.59 1.59 1.40 1.65 1.02 1.18 1.18 1.26 0.82 1.26 0.88 1.26 0.88
4.08 7.76 0.52 0.64 2.24 3.91 1.03 0.73 3.23 3.22 0.78 0.78 3.22 3.88 1.43 1.43 3.22 3.88 6.41	3.21 2.82 2.38 2.7.00 2.72 2.77 2.77 2.77 4.78 4.20 5.05 3.0.10 1.37 1.00 1.10 3.46
AA291137 AA598796 AA447514 R51617 N25344 N40951 R31413 AA156809 AA454562 AA285053 T47625 H24316 AA018215 R00128 R07184 AA706870 AA504845 AA18627 AA18627	AA609731 AA488986 R01515 AA678087 AA405800 R06716 AA496348 W70343 R42736 AA505141 R25403 R39446 AA777555 AA186486 H28091 AA11179 R53943 W86282 AA019335 W86282 AA019335
700503 898098 782576 265668 277185 135065 502527 771323 322786 809503 71312 71312 51850 123061 126829 431462 825742 731290 855236	1031903 743353 824723 123926 430720 742115 126474 755274 345849 31366 825822 133179 23819 448417 628336 162310 624271 40027 415851 36500 731311
4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4	510 511 512 513 514 515 516 517 527 528 529 529 530 531 531 532 533 533 533 533 533 533 533 533 533

Table 2-7: Good vs. Poor Clinical Outcome. Genes upregulated 3 fold or more in at least 1 tumor sample.

4.29 6.75 6.25 6.86 7.75 3.52 1.08 1.13	6.00 6.00 6.00 6.00 6.00 6.00 6.00 6.00	2.0.0 2.0.0 3.0.0 3.0.0 3.0.0 3.0.0 3.0.0 3.0.0 3.0.0 3.0.0 3.0.0 3.0.0 3.0.0 3.0.0 3.0.0 3.0.0 3.0.0 3.0.0 3.0.0
1.30 3.51 0.76 0.96 0.77 0.77 2.35 3.61 2.14	2.0 2.0 2.0 2.0 3.3 3.4 3.4 3.4 3.4 3.4 3.4 3.4 3.4 3.4	2.04 1.00 1.00 1.00 1.10 1.10 1.10 1.10 1
0.24 0.50 1.48 0.70 0.70 0.79 0.32 0.32	6.16 6.16 1.20 0.94 0.81 1.10 0.43 1.76 0.43 1.76	3.24 1.36 1.36 3.20 3.20 3.20 3.35 3.35 3.44 3.44 3.44 3.44 3.44 6.44
0.70 1.02 1.53 1.153 0.67 0.67 3.03	2.55 2.74 2.74 2.19 2.19 2.13 2.13 2.13 2.13 2.13 2.13	0.80 3.63 3.63 2.11 1.17 1.25 0.97 0.95 0.95 0.95
1.78 0.60 3.33 1.15 1.12 1.20 1.20 1.20	2.0 2.0 3.4 4.0 5.0 5.0 6.0 7.0 8.0 8.0 8.0 8.0 8.0 8.0 8.0 8.0 8.0 8	1.23 1.23 1.10 1.10 1.10 1.10 1.10 1.10 1.10 1.1
5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5	8 8 8 8 8 8 8 8 9 9 9 9 9 9 9 9 9 9 9 9	22222222222222222222222222222222222222
13.12 2.10 0.60 0.40 0.58 0.58 26.77 1.39 2.14 1.42	0.73 2.03 2.04 0.09 2.03 2.04 2.04 2.04 2.04	3.54 9.18 9.18 9.18 9.18 9.18 9.19 9.19 9.19
21.82 3.50 0.99 0.67 0.97 44.42 2.31 3.55 3.363	1.25 1.25 1.25 1.47 1.57 1.57 1.57 1.57 1.57 1.57	5.81 6.88 6.88 5.37 2.18 0.19 0.19 1.06 6.73 6.73 6.73 6.73 7.82 7.82 7.82 7.82 7.82 7.82 7.83 7.84 7.84 7.84 7.84 7.84 7.84 7.84 7.84
T59658 R77512 AA459110 AA402760 R38678 R52796 AA872257 AA419542 AA487215 T5280	152840 H08194 AA459293 AA001841 H19327 AA629117 AA629117 AA489219 R36587 N63727 AA644099	AA630445 AA599741 R53951 AA488976 N52837 AA480870 AA486669 AA019320 N59808 AA019320 AA678975 N50843 AA678975 AA788970 AA678975 N50843 AA703219 AA431571 W47000 T80924
76169 145491 814320 727026 23443 41648 1472479 245624 490647 769796 841308	68061 45391 810301 427396 50926 1035765 47404 825060 137275 293539 845380	854633 1070062 40136 810358 824720 244391 814547 840990 363055 288959 1048789 1240220 383528 280950 825669 784216 435992 1636251 78246 325511 109316
533 534 536 537 540 542 542 543 543	544 545 547 553 553 553 553	558 558 559 560 561 563 563 563 563 574 575 575 575

Table 2-7: Good vs. Poor Clinical Outcome. Genes upregulated 3 fold or more in at least 1 tumor sample.

0.84 1.11 0.78 0.78 1.26 1.26 1.35 0.79 0.79	0.78 0.78 0.99 0.99 0.78 0.79	7.16 0.38 1.03 1.03 1.03 1.03 1.04 1.04 1.05 1.05 1.05 1.05 1.05 1.05 1.05 1.05
1.23 1.06 3.08 0.87 0.89 1.44 1.30 1.30 1.11 1.70 0.74	2.0 4.4.1 1.0 1.0 1.0 1.0 1.0 1.0 1.0 1.0 1.0 1	0.95 1.07 1.07 0.99 0.90 0.90 0.90 0.90 0.90 0.90 0
1.05 1.10 2.21 2.21 1.05 3.05 3.05 0.19 0.30 0.30 3.47 3.54	3.18 3.80 3.80 0.65 0.05 0.07 0.22 1.68 1.68	3.42 3.21 1.03 0.63 0.77 0.33 0.50 1.20 1.20 1.20
1.01 1.35 1.28 1.28 1.15 1.06 1.08 1.09 1.09 1.09	2.82 2.82 2.05 1.05 1.28 1.28 1.09 1.13 1.63	1.72 0.60 0.60 0.73 0.73 1.28 0.98 1.10 0.67 0.67 0.67 0.67
3.89 2.1.54 2.1.10 1.10 1.10 1.10 1.10 1.10 1.10 1.	1.02 0.52 1.16 3.17 3.56 0.82 3.97 1.05 0.38	0.13 1.36 0.60 3.72 3.72 1.92 1.13 1.06 3.52 0.12 3.52 0.73
09:1-1-1-1-1-1-1-1-1-1-1-1-1-1-1-1-1-1-1-	2 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5	1.57 1.57 1.56 1.56 1.56 1.56 1.56 1.55 1.55 1.55
0.00 6.00 6.00 6.00 6.00 6.00 6.00 6.00	0.25 0.25 0.27 0.37 0.32 0.34 0.34 0.34 0.35 0.35 0.35 0.35 0.35 0.35 0.35 0.35	0.54 6.24 7.47 7.47 7.48 0.55 0.69 0.27 2.70 0.27 0.27
0.62 0.86 0.88 3.44 1.24 1.74 0.82 2.88 5.53	2.13 0.45 1.56 1.14 1.14 1.75 1.75 2.91	0.84 9.80 0.74 2.30 2.47 2.4.12 0.85 1.10 1.10 1.10 1.10 1.10 1.10 1.10 1.1
H29268 T99011 AA521431 R63106 R37519 AA479967 N21338 AA001449 N24538 AA411686 T55728 AA434092	AA677534 AA021628 R52901 R24506 H74265 AA443832 AA663123 AA663123 AA663123 AA639223 AA504265 AA399223	N58170 AA454634 AA214031 W60845 AA677287 AA705319 N47240 AA282134 AA282134 AA505051 AA778653 H23202 R12808 AA035171 R38017 AA620715 R44447 AA620715 R44447
49842 122684 826173 137940 137254 753982 265103 361974 267541 753378 73638 624490 377731	460403 364083 41569 131791 133273 229365 784065 852548 24855 196387 825416 725745	247660 811880 562395 341805 454771 462856 280459 711918 825654 1049009 52079 52079 25384 471892 24131 1049230 34869 431505 250988
578 580 581 581 582 583 584 586 586 589 589	592 593 595 596 600 601 603 604	605 606 607 607 611 611 612 613 614 619 619 620

Table 2-7: Good vs. Poor Clinical Outcome. Genes upregulated 3 fold or more in at least 1 tumor sample.

	•	_		_	•	- •	- •	- ,	,	- `	_ `		•			•	•	•	•	•	•	•	•		•		•																				
26.0	1.10	0.93	- 5	77.	7 · ·	5.5	02.1		1.01	9.0 48.0	1.28	0.76	0. 26.	0.98	0.93	3.85	0.67	0.91	0.95	0.33	0.77	1.20	0.89	0.93	1.7	3.55	1.1 0	0.76	1,12	1.47	3.30	1.07	1,13	94.0 0	0.80	27.0	† *	70 +	7. 0	8	9.0	4 60	9 90	7.70	-		
7 20	5 5	. 40	7 60	17.1	וני.	1.73	1.47	1.24	1.16	1.06	1.47	1.37	1.38	0.87	1.06	1.18	1.30	0.86	0.86	0.42	1,33	0.93	0.77	1.29	1.03	1.38	1.17	0.56	. .	0.68	0.51	99.0	0.57	1.24	0.95	0.70	5.05	40.0 40.0	98.0	1.6.1	7.52	94.1	1.32	1.20	0.96		
Ç	3.23	2.55	4 .06	0.81	1.13	0.43	0.68	1.10	3.39	3.19	3.62	3.01	0.88	0.86	3.27	0.45	1	. 6. 6.	0.42	1.25	<u> </u>	37		1.43	0.39	38	388	1.27	3.16	0.75	1.96	0.67	0.88	0.26	3.44	0.68	0.71	0.58	3.07	3.20	3.06	3.96	3.58	0.81	0.76		
	1.21	0.95	0.89	1.27	0.90	1.1	0.53	1.24	į -		22.0		5 +	1.10		. c.	7 7 7) 	9 C	- C-			9 9	7.0	* i	67.	6 6 6	7.00	1.1.1		2.5	0.82	0.93	0.95	1.82	0.80	0.58	0.71	1.39	0.79	0.94	0.69	0.99	0.81	1.17		
	1.12	0.83	0.89	3.14	3.07	1 12	- 6	 	5 5	- 6	90.0 93.0	, c	12.7	3.35	9 9	1.53	0.93	3.16	1.26	4.69	1.44	3.42	0.54	3.72	3.20	3.15	0.67	0.63	3.22	0.52	7.0	0.08 8.33	, k	 56.	4	3.61	0.48	4,08	0.82	1.10	66.0	0.69	0.82	3.40	3.47	: ¥	2
	1.55	1.54	154		5 7	<u>.</u>	đ.	\$ i	4. 4.	. .	1.53	1.53	1.53	1.53	1.53	1.53	1.53	1.52	1.52	1.52	1.52	1.52	1.52	1.52	1.52	1.52	1.52	1.52	1.52	1.51		1.51		 	 	. <u>.</u>	. .	 	. t	5 &	5.4	5.5	5 4	5. 5. 5. 5.		5.	
	0.80	7	5 5	17.4	94.O	0.23	2.16	2.14	0.19	5.62	0.69	75.80	0.55	0.71	0.33	1.11	10.13	0.56	0.76	7.5	2.36	0.62	9.11	0.88	2.12	2.53	4.09	24.63	0.50	8.15	1.20	2.73	0.18	0.47	(4.81 18.43	Z.Z.3	0.70	C.53		78.	- c	 	5.21	4.0	07.1	0.71	
	4,23	9 6	S 6	54.	0.75	0.36	3.32	3.30	0.29	8.63	1.07	116.27	0.85	1.09	0.51	1.70	15.46	0.85	1.15	2.35	3.59	9.9	13.85	7.	3.23	3.84	6.20	37.32	0.76	12.33	1.81	4.13	0.28	0.71	52.58	6.40 0 1	7.7	0.53	2.63	2.68	2.20	21.1	8. i	0.71	1.79	1.06	
	A A FORROO	7230000	780047	AA489016	AA135886	H15695	AA040617	AA879437	R51912	N51335	AA418414	N52178	N75356	H72959	AA015793	N26724	AA142922	R82733	AA196465	AA872397	R10378	R33103	AA701502	AA129552	H57857	H02307	AA290737	N48809	R16245	AA159578	T60149	AA733177	AA284113	R40176	174105	AA424562	N93428	N48580	AA773810	AAZ78320	163031	AA707008	W46985	AA421284	W31725	R44985	
	00000	89/820	165828	824920	502674	49203	486208	1505692	39593	283142	767284	284247	298862	213917	360478	266300	505225	149742	A28295	1472743	120024	135800	435470	564803	205497	151201	713922	279481	53393	592594	81408	399421	324751	27769	84613	767176	307231	279249	878277	703808	80772	451351	325088	731051	320797	34442	
		623	624	625	979	627	628	629	83	631	633	633	3 2	535	35,0	23.6	3 6	3 8	3 8	3	3	3 2	} }	£	7	3	3	3	650	651	652	653	654	655	656	857	658	629	99 99	8	662	663	8	665	999	299	

Table 2-7: Good vs. Poor Clinical Outcome. Genes upregulated 3 fold or more in at least 1 tumor sample.

- 1	- •	- •	•	•	_ •	•	- •	•		·																																		
4.43	0.82	0.83	05.7	7.0	7.1	8.5		81.T	0.80	0.7g	0 6	7 7 7	70.	2.6	. 6	9.0	5,5	7.00	3 3	5 6	 	0.55 50.0	3.25	3.26	0.73	0.61	\$;	0.98	08:0	1.15	0 C		5.5 14.5	0.87	0.72	1.06	0.92	20.0	9.56	6 6	0.32	 	20:5	
0.76 0.27	1.75	1.35	1.72	0.85	3.20	0.46	0.63	2.09	1.07	1.56	0.93	1.23	1.76	9.6	0.00	re.0	3.23	69:0	1.32	0.97	0.74	1.28	0.98	0.95	0.81	0.78	0.73	96.0	1.01	0.86	1.17	0.67	5 C		o	2. 4 40 4	0.9	0.00	0.0	2.03	1.37	1.72	0.71	
0.50	3.40	3.49	3.49	3.53	1.15	0.42	0.52	3.00	0.95	3.09	0.98	0.91	ы 9	3.34	0.97	3.63	1.59	2.15	3.16	3.27	0.35	3.63	0.80	0.74	3.88	4.22	9.64	0.87	0.72	4.6	0.88	0.51	3.91	0.83	3.33	3.23	3.73	3.01	3.93	3.14	0.89	3.04	3.87	
1.01 81.8	20.5	1.25	0.58	50.	1.31	3.64	1.45	0.95	1.01	1.45	0.97	0.93	0.73	0.92	1.06	0.91	0.78	3.02	1.05	1.02	1.49	0.97	98.0	1.15	0.79	0.81	1 12	0.88	1.45	1.19	1.05	1.33	0.65	- \$.	1.35	0.86	0.82	1.28	0.67	0.69	1.06	0.64	1.08	
77.0	0.38		0.33	, ,	14.0	9 6	, c.	3.50 0.50	0.40 7.40	75.0	 	3.07	0.56	1.13	15.	5	0.77	0.85	000			96	4 6.34 4 8.4	. t	1.4 1.4	± 00			2.00	3.55	. C	3.96	0.62	0.69	0.74	1.12	0.58	1.39	0.65	0.69	3.01	1.06	0.94	16
1.49	1.49	1.49	1.49 0.49	D (*)	1.49	1,49	1.49	1.49	 64.	04:- 04:-	94. 64.	04.L	2 7	. τ δ α	. 4	4.4	 	· ·	9.40	7.4.	7.4.	7.47		7.47	1.47	1.47	1.46	1.46	1.46	7.40 0.40	04.1	04. 4.	. t.	1.45	1.45	1.45	1.45	1.45	1.45	54. 74.	 	 - 4		} •
3.07	0.78	1.01	7. 20.	10.74	4.0	1.16	4.4	0.82	6.83	0.53	0.41	4.56	ر د ز	15.21 0.00	0.0 0.0	\$ (S)	6.02	0.0	1.35	1.25	0.80	0.46	1.40	6,52	5.09	0.46	0.79	0.59	0.49	0.70	0.39	0.63		£4:01) () ()	6.5	3 4	34.17		777	5.5	T.S.1	7.46	co.r
4.59	1.17	1.51	2.45	16.03	0.65	1.73	2.15	1.23	10.16	0.78	0.60	6.76	60.	22.54	9. F	0.50	8 8 9	0.74	1.99	2 .	1.18	0.68	5.06	9.58	7.47	0.67	1.15	98.0	0.72	1.33	0.57	0.92	0.22	\$ 5 \$ c	0.45) () ()	20.0	0.4.0	9 6 9 6	χ. Σ. (1.52	9.5	10.78	1.5.1
W16832	AA598817	N35888	W87826	AA703161	AA454928	R08121	AA873578	AA131406	R26785	N33443	R44544	AA400457	AA609392	AA702077	R12385	H03436	N69528	AA489729	AA017468	AA962119	AA701900	N58417	AA707400	H08899	N30285	AA644211	AA678203	AA283949	W86214	R26444	H19320	AA405569	AA453085	10407	100/21	K32320	AA/01046	H02039	AAZ13009	R93591	R66082	AA233070	AA488064	R26206
320392	897956	272529	417081	435577	814739	127120	1472152	503617	132636	270750	34007	753162	743481	384224	128165	150897	294503	823663	361291	1600281	435536	248117	451546	44975	257504	845477	430864	700571	415868	133150	50918	772425	789069	219888	109863	38977	397432	208377	683278	197731	140174	666492	840686	133053
8	3 8	670	67.1	672	673	674	675	678	877	678	679	089	681	682	683	8 8	685	989	687	889	689	069	69	692	693	95	895	989	269	869	669	8	5	702	5	ğ	202	8	707	% %	60 2	720	711	712

3.19 3.19 3.19 3.05	0.846 0.92 0.92 1.25 1.22 1.23 0.43 1.33 1.33
1.61 2.98 2.98 0.91 0.68 1.00 1.10 0.86 0.83 0.67 1.13 0.67 0.94 0.98 0.98 0.98 0.98 0.98 0.98 0.98 0.98	0.83 0.83 0.82 1.12 1.09 0.38 0.95 1.19 0.95 3.08
3.63 3.63 3.63 3.03 3.03 3.03 3.03 3.03	0.71 3.26 3.38 3.32 3.32 3.34 0.64
1.10 0.49 0.87 1.38 1.29 0.90 0.90 0.90 0.92 1.13 0.92 0.92 0.93 0.94 0.93 0.94 0.95	0.53 1.21 1.04 1.04 1.04 0.64 0.90 0.90 0.92
0.47 1.155 1.155 1.18 3.00 3.26 3.63 3.63 3.63 3.19 0.65 1.14 1.24 1.26 0.09 1.28 3.28 3.28 3.28 3.19 0.57 1.20 0.65 0.65 0.65 0.65 0.65 0.70	1.45 1.28 1.01 0.79 0.23 0.56 0.59 0.59 0.54 1.03
44444444666666666666666666666666666	4444444444444444444
3.41 72.70 1.08 1.08 1.08 0.22 1.28 0.48 0.48 0.49 1.20 1.20 0.39 0.39 0.39 0.39	2.44 1.40 1.07 1.07 1.07 2.13 5.29 10.48 33.92 0.14
4.93 104.94 1.56 62.25 69.25 0.69 0.37 10.24 1.18 1.18 1.10 1.10 1.10 3.20 1.30 1.30 1.30 1.30	5.00 6.70 1.97 3.44 1.50 25.61 3.88 58.28 7.43 14.69 0.20 4.21 47.45
N67766 AA872383 AA455659 AA683050 R47979 N63286 W90067 T97890 T90980 AA974971 R39066 AA974471 AA625890 AA677457 R25823 R437241 H87106 H87106 H87106 R75338 AA704332 AA668470 R42533 AA704332 AA48599 AA707121	AA480906 N51740 H60696 T62040 N58052 R07196 AA432124 AA432124 AA432124 AA481052 H62529 H92216 AA460282
291426 1472735 814018 971367 153411 290162 418081 121530 143759 489800 112482 1555523 24915 757451 239446 392607 744944 460218 39393 28393 28379 261851 32495 33269 332706 853809 30275 452059	713193 82734 814595 281737 208985 85609 24746 126847 781467 451918 269563 814661 208165 221778
7 4 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5	743 744 748 749 750 751 751 751 751

Table 2-7: Good vs. Poor Clinical Outcome. Genes upregulated 3 fold or more in at least 1 tumor sample.

Ψ,	_	-	•-	-	-	•	-	*	-	-	•	• •	- •	- •	- ,	- ,	_	-	-	-	•	-	•		- •	- •	- •	- ,	- ,	- ,	, ,	-	•	••	Ψ,	- ,	- •	- •	- •	- •	- ,	- ,	-	-	•-	-		
0.81	1.24	1.14	0.78	1.05	1.01	1.12	1.08	0.75	990	. t	5 4	5 .	17.1	8°.	01.1	0.48	. 8	0.78	1.58	1.10	3.68	0.72	1 a	3 6	- 6	0.0	97.1	3.90	0.70	0.73	. 8	0.83	0.98	0.64 4	1.28	1.14	0.86	0.3	04.1	3.18	24.0 54.0	0.47	4.09	0.8 48	0.54	1.01		
0.71	1.27	28.0	0.80	0.80	0.97	108	0.63	5.6	5 6	67.0	D (C	0.50	0.78	0.39	1.25	3.57	0.77	0.68	9.0	98	5 5	- a	9.6	S 6.0	1.22	1.09	0.0 40.0	0.65	0.88	1.77	0.39	0.70	4.06	1.15	0.93	1.24	0.47	0.89	0.31	1.39	0.30	0.57	0.87	0.60	1.73	0.99		
3.16	3.27	3.62	3.10	2.47	, c	20.0	7 7	4 6 6	0.52	3.07	3.29	0.69	3.20	4.28	0.24	1.22	3.68	06.0	90 8) -	÷ 5	- G	1.63	3.92	3.07	3.26	3.06	0.35	0.82	3.09	1.74	1.13	0.38	3.03	3.35	3.07	0.57	0.53	3.38	3.07	4.54	3.54	3.19	0.91	3.16	3.07	;	
1.56	0.50	200) c	2 0	0.0	0.80	1.0.1	1.18	0.95	9. 28.	1.18	1.10	1.40	0.79	0.53	0.83	9 6 7	2 6	28.0	6/.0	1.31	3.06	0.84 48.	1.03	0.79	0.82	0.86	0.88	0.95	3.17	3.37	0.74	7.70	0.71	0.78	1.27	1.06	0.97	0.64	0.86	1.71	0.86	0.85	22.0		3.	
0.75	9 0	8 6	3 6	90.0	0.73	0.63	0.40 84.0	0.97	3.88	0.95	0.97	0.49	0.62	0.45	3.54	1 10		5 6	10.0	99.0	0.57	0.85	0.36	0.48	0.80	0.83	0.72	3.05	3.53	0.26	55.0	4 6	0.0	, c	27.0	0.54	3.60	1.11	0.69	0.46	0.59	0.43	7.		÷ 6	9 5	o !	8
4	£ .	 9. :	1.39	1.39	1.39	1.39	1.39	1.39	1.39	1.38	1.38	1.38	1.38	38	- + & &		000	35.	3.38	1.37	1.37	1.37	1.37	1.37	1.37	1.37	137	+ 36. +	t	5.4	 9. 4	o	٥٠. ٥٠.	05.1 05.1	0.5 35		35.	35	1.35	1.35	25	- - -	<u> </u>	5 6	\$ 3	\$ 7	4.7	
9	2 :	15.21	19.57	0.00	1.32	34.84	52.40	1.03	0.42	1.01	2.08	2.42	i C	20.5	. ·	87 CO C	207	0.73	0.52	0.57	102.66	0.10	13.04	1.68	0.20	 	- 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4	5.5	<u> </u>	7.97	79.7	26.33	2.60	1.37	2.32	2 6. 10	900	5.5	. 6	51.24	78.0	. v		0.70	1.49	0.69	0.37	
ć	29.0	21.24	27.27	0.13	2 .	48.51	72.84	1.43	0.59	1.40	288	, i	7 7	1.1	9 9	A/:	2.79	<u>-</u>	0.71	0.78	140.80	0.13	17.85	200	90	5.50	10.4	\$. ?	4.7	67.1	49.01	35.81	3.53	1.86	8.75 5.25	70.81 70.43	33.15 00.0			200	20.00	- u	6 6 1	5.73	2.00	0.92	0.49	
	1092X	H84759	AA702193	AA054439	AA598775	H81716	H65300	H04757	R17747	AA678361	R56840	R00707	AAEBOKEE	NO COL	14603U	74405/	AA43036/	AA430410	R19878	AA704749	H94870	AA776702	N62394	AA582642	A4431772	D70508	A A 2 7 0 1 7 2	44000000	A44246	A445110	AA2919/2	AA425692	AA447764	AA434403	AA428341	132213 A A 45 4036	A4733540	NISEADO	H86276	AA708348	A 8 9 5 7 4 0 6	7007490	011047	8/08/1	AA253442	H92875	W44938	
	144932	219937	384134	381054	898058	219861	208225	152270	2539A	431866	42826E	130433	47474	451268	309081	33854	769857	769947	34888	451161	230236	1292829	28863	450680	100000	105301	141004	7040/0	1048696	809488	725364	773204	813661	770879	769673	9//122	614/44	02000	22224	200673	392073	1304031	01066	248734	669367	231718	322857	
į	758	759	760	761	762	763	764	765	788	787	2 4	8 8	ê î	e i	5	77	13	774	775	776	111	4	3 2	700	3 5	5 6	7 67	3	\$	785	38	787	788	789	8	F6/	78/	2 7	5 5	2 4	8 5	2	86/	789	8	<u>8</u>	802	

Table 2-7: Good vs. Poor Clinical Outcome. Genes upregulated 3 fold or more in at least 1 tumor sample.

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0.33	1.20	1.85	1.05	3.01	4.7	0.65	3.11	0.70	0.96	0.59	0.66	3.43	0.88	0.72	0.83	1.48	0.95	0.91	0.84	1.13	0.55	1.02	1,15	1.13	1.03	0.80	0.82	0.98	0.60	0.87	1.16	o (.	 8	0.43	0.87	1.20	1.15	0.69	0.62	0.85	1.08	0.70	0.68		
0.58	1.47	0.67	200	9 5	3 5	0.51	1.02	3.43	1.10	0.55	0.82	0.77	1.13	0.95	1.05	0.97	0.45	1.19	0.85	0.76	1,33	0.73	1.01	0.95	0.96	0.62	0.99	0.61	1.18	1.17	1.02	0.68	5. 6. 6. 6. 6. 6. 6. 6. 6. 6. 6. 6. 6. 6.	0.00	9 6	278	88.0	0.41	103	0.76	0.87	97	98.0	}	
4 83	50.6	5 6	t C	0.00) Y	 	3 6 6	26.0	0.9	5.50	980	0.00	9 6 40	3.63	3.40	0.50	55.0	3.16	3.49	600	9 6	9 8		3 5	3.25	0.62	8 8	0.67	3.40	3.01	3.00	3.29	0.21	3.46		5.32	6.5	5.00 P 8 7	9 6	0.00	5.52 ac	3 6	9 9	6.0	
į	C:0	0.52	3.32	19.0	0.92	9.0	00:0	6.0	8 6	. T	t 6	9 9	2 6	2.0	500	2 6	* °	9.50	t 6	2 5	0.73	5.5	0.0	60.0	C.93	9.0	, c	1.14	0.61	0.62	0.63	69'0	1.39	0.86	80°C	0.74	0.66	0.62) i	o 6	0.58 0.98	0.00	\$! - 0	3.6/	
	0.35	0.46	0.52	0.67	40.	0.41	1.19	0.85	0.80	3.14	4.38 6.60	0.42	1.18	0.7	- 3	0.00	3.73	0.23	[9.0 [9.0	0.63	3.68	0.68	ы 9	0.63	3.74	0.0	5.70 2.70	. c	3.02	22.0	09.0	0.99	3.10	0.51	0.51	0.50	3.13	0.61	0.55	0.59	0.79	0.68	0.50	0.62	19
	1.33	1.33	1.33	1.33	1.33	1.32	1.32	1.32	1.32	1.31	1.31	1.31	1.31	1.31	1.31	1.31	1:31	1.30	1.30	1.30	1.30	1.30	1.30	4.30	1.29	1.29	1.29	1.29	87.1	2.50	1.20	1.28	1.27	1.27	1.27	1.27	1.27	1.27	1.26	1.26	1.26	1.26	1.26	1.25	
	2.54	78.95	1.93	26.74	0.52	4.55	0.10	0.38	0.63	2.29	11.10	9.64	1.12	0.50	5.91	11.97	0.3 4	11.49	3.82	10.08	4.38	1.12	2.21	37.28	2.06	9.31	0.93	0.86	1.09	ct.:	94.0 C	0.65	2.08	0.21	2.88	0.32	9.82	23.43	2.13	1.28	2.16	17.73	0.80	1.72	
	3.38	105.05	2.57	35.54	0.70	6.02	0.13	0.51	0.83	3.01	14.58	0.84	1.46	0.65	7.73	15.65	0.45	14.99	4.99	13.13	5.71	1.46	2.86	48.29	2.66	12.02	1.19	1.1	1.40	1.47	4.47		2.66	0.26	3.67	0.40	12.44	29.64	2.68	1.60	2.72	22.29	1.01	2.15	
	AA676920	W37532	AA497111	H66675	H09172	AA682838	AA758470	AA432134	AA447592	AA417921	AA873159	AA401826	AA464979	AA504505	AA976561	AA704908	AA927544	AA485303	AA677337	H69553	T63324	AA427561	R00283	N62763	H11732	AA701328	AA164836	AA778663	H15549	AA481437	AA/046/0	760503	AA279060	AA485249	N47443	AA774501	AA157813	R66923	AA677300	AA482127	N50859	N76088	AA486765	N59532	
	460164	322136	823562	211387	46518	450300	431029	781492	782687	767405	1472689	725224	810109	825366	1558655	461363	1552618	840266	454469	212473	80109	770059	123278	289562	47481	435126	595148	1049030	49499	756554	383752	645079	703046	815835	280697	399264	588915	140299	454795	756450	280982	295454	841057	248631	
	803	2	S S	8 8	3 2	808	808	810	811	812	. 813	814	815	816	817	818	819	820	821	822	823	824	825	826	827	828	828	830	83	832	833	83.4 4.5	3 8	3 6	88	839	840	24	842	843	4	845	846	24	

Table 2-7: Good vs. Poor Clinical Outcome. Genes upregulated 3 fold or more in at least 1 tumor sample.

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94.0	9 ; c	- u	96.0	9.0	0.00	2.6	3.63	7 7	(O.O.	9 5	0.0	20.4		5.5) e	1.8	1.10	3.83	0.82	0.78	0.75	7.0	0.82	0.83	3.05	75	0.68	2	0.69	1.25	0.20	4.0	0.58	0.32	0.23	0.52	0.00	0.59	0.72	3.18	3.09	0.62	3.03	•	
•	0.40	5.0	49.0	1.05	88.0	D .	4 6 6	0.27	0.79	D	0.5	0.00	1.85	LC:0	0. C	5.0	220	200	0.90	3 6	- 88	0.00	0.0	8000	0.00	0.52	48.0	0.78	0.63	0.08	0.22	0.65	0.21	0.68	0.55	0.55	0.00	0.14	3.02	20	0.45		0.62	;	
•	0.34 5	3.03	3.25	3.20	3.28	0.40	0.70	0.63	0.71	0.72	0.29	0. 20.	1.20	0.17	3.20	3.03 9.03	0.0	† ;	\$ 6 6 7	20.0	3.30 0.00	10%	3.14 4.00	0.20	15.5	0 0 0 1 0	3 40				 	3.26	0.24	3.29	0.30	3.25	1.88	0.37	0.54		3.0	200	3 c	- - -	
•	0.92	99.0	0.68	09.0	0.64	0.84	0.63	0.89	0.88	96.0	1.71	<u>.</u> 4	3.16	1.07	1.12	0.80	0.59	0.00	0.74	0.74	0.74	0.84	1.04	0.63	0.66	0.86	4.0	\$ C	C:0.	20.0		 98.0	0.73	1.12	105	107	3.69	202	 	3 6	7 C	C:0	0.00	t	
	3.84	1.04	0.41	0.79	0.56	3.33	0.52	1.32	3.17	3.05	0,49	0.21	0.00	3.14	0.40	0.63	3.10	0.53	0.24	0.67	0.49	0.63	0.54	0.52	0.60	, , ;	3.12	0.62	0.51	3.33	3.28	87.0	600	9 6	0.0 50.0	0.01	3 6	9.00	0.4 0.4) ; ; ;	9. 0 4. 1	0.73	3.01	0.77	20
	1.25	1.25	1.25	1.25	1.25	1.25	1.25	1.24	1.24	1.24	1.24	1.24	1.24	1.24	1.24	1.24	1.23	1.23	1.23	1.22	1.22	1.22	1.22	1.22	1.22	1.20	1.19	1.19	1.19	1.16	1.15	61.13	 	 	2 7	<u> </u>	7.17	= ;	1.10	1.08	1.07	1.03	1.03	0.97	
	4.19	1.47	60.5	2.25	15.13	2.25	18.72	3.79	1.52	35	x 47	: e	800	90.0	2.89	2.45	11.12	17.78	4.32	9.45	3.92	0.58	4.94	14.12	31.16	5.54	8.11	2.96	6.54	0.95	15.23	3.68	67.7	13.05	7.9S	05.5	30.90	0.02	0.02	2.14	7.59	6.99	10.12	17.01	
	5.25	184	72.8	28.0	18 84	2.81	23.3	4 7.1	80	. t	5.4	; ;	: 2 : c	0.0	3.57	3.02	13.72	21.84	5.30	11.57	4.80	0.71	6.03	17.18	37.89	6.64	9.69	3.53	7.79	1.10	17.56	4.23	4.	15.02	5.0	د. مي	84.58	0.03	0.02	2.31	8.14	7.22	10.45	10.70	
	R34225	AA491261	AA707080	AA213667	HA6559	AA451863	R92281	A A 4 5 4 9 8 6	AA465194	HEA364	A A DO 1 4 4 4	N32074	AA845584	AA425298	AA701976	AA676865	H29500	N29457	R55965	H13438	AA757711	AA703159	H38848	AA504132	N21688	AI015196	R34273	AA704278	AA699719	R16146	AA465495	R01304	H10072	AA4/9284	AA623979	404407	163245	AA057195	AA131885	AA521384	N49107	AA625666	AA478470	N91767	
	136028	02020	45450	401080	222480	706308	106180	944044	4 000	01000	203330	50100	200100 440400£	773284	435953	460143	52730	270331	40636	148836	396085	435506	190692	825234	266455	1623328	136070	450997	433300	53158	814053	123755	46843	754280	745490	141361	80050	489076	504358	826995	279800	745347	786602	306743	
	RAR	2 2	9 6	9	- 63	700	000) H	000	2 6	200	0 0) () () ()	200	282	853	200	865	998	867	88	840	870	871	872	873	874	875	876	877	878	878	880	881	882	883	8	885	886	887	888	889	830	891	

Table 2-8: Good vs. Poor Clinical Outcome! Genes upregulated 3 fold or more in at least 2 tumor samples.

Table 2-8: Good vs Poor Clinical outcome: Genes upregulated 3 fold or more in at least 2 tumor samples

ndn	gulated	upregulated 3 told of more in account							VA/06	Fold(mpm35/AV	# of Tumors
	IMAGE	Gen Bank			Fold(AveGood/Av Fold(mpm27/Av Fold(mpm28/Av Fold(mpm29/AV Fold(mpm29/AV Fold(mpm27/Av Fold(mpm27/Av Fold(mpm28/Av Fold(mpm29/AV	old(mpm27/Av l	old(mpm28/Av	Fold(mpm29/AV	ePoor)	ePoor)	Samples
Order	Clone ID	Accession	Avecood	AVELOG	ePoor)	ePoor)	eroor)		000	7.30	4
		Legun	1	4	70.3	7 07	6.80	3.51	0.0) c	4
-	299679	W05628	2.55	0.50	2.0	5.0	3.61	6.33	4.17	07.0	rø
7	811162	AA486471	28.87	6.83	4.41	\$ 5	2.20	4.22	2.10	C1.12	,
e,	471641	AA034945	11.03	1.08	10.17	20.03	40	0.27	9.85	7.08	~ (
4	281039	N47717	36.67	4.42	8.29	75.27		0.67	7.05	8.80	
140	282089	N48259	27.13	3.45	7.87	22.41	- t	130	9.52	15.90	m
• •	155072	R71393	3.32	0.45	7.44	8.50) (a)	8 6	5.30	10.17	က
1 0	208210	H65286	2.70	0.42	6.48	14.00	9 6		8.30	5.96	က
. α	770066	AA430545	26.72	4.62	5.78	12.05	, c.o.	- C	4.37	12.92	က
9 0	712950	AA282273	38.22	6.94	5.50	8.8 4	2 8	6 6 6	8.59	3.28	ന
• •	81331	T60111	4.13	0.82	5.03	11.18	J. C	999	3.86	5.42	က
2 =	287728	N59158	3.49	0.71	4.90	14.39	 	4 46	2.14	3.71	က
: \$	41228	R58953	0.05	0.01	4.38	9.07	2.03	3 t C	000	3.18	က
<u>.</u>	122345	T99191	0.03	0.01	4.16	12.66	÷ •		4.30	9.31	ო
2 7	186918	H43317	6.27	1.60	3.93	3.28	- c 4 c - c	54	2.55	5.03	က
. A	1049033	•	2.50	0.65	3.87	4.86	7.30 7.30	88	3.51	6.23	ო
, ,	195365		1.81	0.50	3.62	6.03	4. 5.	5 4	2.28	0.54	က
<u></u>	50114	H16743	1.48	0.43	3.43	5.36	4. 0. c.		1.32	1.35	ო
= =	838611	_	87.41	25.99	3.36	5.71	77.4	1.5	3.24	3.77	က
2 4	283198		5.16	1.55	3.32	7.31	, C. O.	, 2	4.76	3.74	ო
2 5	745525	_	4.16	1.27	3.28	5.86	1.7.	5 6	1.59	3.11	ო
7 1	159487		0.12	9.0	2.92	2.50	, c	2 2 2	0.79	3.97	ന
3 :	1034473	•	76.12	27.06	2.81	2.22	3.21	135	3,32	3.28	က
1 %	138444		2.02	0.74	2.75	4.16	- •	- c	3.13	3.31	ო
3 %	378488	AA777187	17.01	7.17	2.37	3.23	02.1	28.00	1.49	1.02	7
3	767130	AA424535		0.37	11.65	1.60	20.03 7. 4.7	44 45	1,65	1.22	7
8	307337	N95226		0.96	11.29	1.65	† †	33.58	9.	3.17	8
27	502625	AA134576		0.90	8.08	4.6	u	15.74	05,1	0.71	8
88	41869	R66438	1.41	0.28	5.03	1.46	0.0 7.0 7.0	2.93	0.99	7.09	6
83	301735		4.10	0.90	9.56	9.03	 	11.13	1.94 46.	2.65	7
8	136605		32.22	7.39	4.36	7.55 6.45	38	0.78	6.56	7.23	74 (
ಕ	362278	•	5.53	 1.4.	3.91	40.0	4 50	10.09	2.99	<u>.</u>	71 (
33	756708	⋖	2.58	0.67	8. 20.		5. 4 36.	11.96	0.59	0.98	7 (
g	265028		1.22	0.32	3.78	- c	9 8	11.75	0.59	0.31	7
z	1409509	_	2.84	0.75	3.78	0.30) () ()	10.11	1.36	3.00	7
8	365707	•	<u>+</u> 2	0.53	3.68	0.30 0.00		10.03	0.93	0.97	7
8	768953	٩	3.43	0.9 4	3.66	0.5 0.0	- K	990	2.00	9.41	7
37	357628		1.80	0.49	3.65	2.67		207	<u>*</u>	1.38	7
8	35058	R45192	28.75	7.90	3.64	9.1.6 7.1.6	1	i			
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N	.4	.4	•	,,	"		. (• •	• `	•	. •	••	••	••	••	•	•	• •	•	•	- •	- •				. •	- •	•	•													-								
9.44	8.17	4.57	2.75	0.88	9 80	7 60	200.7	0.30	すべつ	3.69	3.25	9.18	1.08	8.89	0.75	0 10	3.16	6/:1	4.74	5.7	2.19	1.24	1.47	7 80	4.05 4.44	; c	4	0.72	1.24	0.76	4.89	2.27	3.42	6.33	1.06	1.03	3.11	4.38	0.61	0.63	3.26	1.99	40.4	1.12	2.25	. d	9.00	5.00		
5.51	3.97	000	27.6	7.7	n t	2.55	1.60	96.0	2.50	2.35	2.42	1.32	9	9 6	n c	3.20	0.0	1.70	6.81	4.53	1.58	28.0	9 6	7 5	7.1	6.48	2.55	0.46	2.85	0.69	1.56	3.01	2.18	4.00	1 .9	3.15	3.69	5.65	1.27	1.21	2.49	1 76	. u	ţ °	2 5	2.5 20.0	0.28	2.42		
27.0		7.00	Z.30	90.6	10.28	0.51	1.61	2.26	1.26	0.59	5	9.0		6 ! 6 !	1.77	0.57	0.74	4.46	9.0	1 52	. c	o :	6.22	1.02	0.79	1.14	1.55	7.31	4.22	7.85	1.62	17.1	5.5	0.65	6.17	5.37	2.58	1.09	5.67	, r	3 5	7 7	6.9	[6.0 [6.0	5.61	1.56	0.22	0.76		
ç e	0.52	2.44	1.91	2.54	3.45	1.09	4.91	3.72	2 96	1.26	07.7	0.40	0.81	5.84	0.71	0.89	4.10	2 99	1.13	2 7	1.4.	6.21	3.75	3.46	1.98	0.96	1.25	4.28	20.0	25.0	7 2	777:- 0 0 0	4.00 4.00	- c	. c. 4	- c	9 6	07.7	9 6	n 7		3.72	3.10	0.53	3.38	2.03	3.78	1.42		
	1.47	1.63	1.01	3.72	0.86	8.38	9 6	60 a	9 6	9.0	8.02	7.62	3.86	6.21	3.68	10.04	0.74	200	D 10.00	0.40	0.82	0.92	1.83	6.94	4.81	85.0	6.50	69.0		3.0.	3.32	3.79	3.41	5,42	1.63	3.54	7.57	2.17	44.0	1.63	0.43	0.61	3.24	0.31	0.80	3.47	0.70	3.36	2	
	3.53	3.51	3.36	3.35	200	9.00	5.63	3.28	3.20	3.23	3.18	3.14	3.12	3.12	3 1 1	000	5 c	9 6	2.86	2.81	2.81	2.81	2.77	2.75	2 6	. i.	2.72	2.71	2.68	2.67	2.63	2.62	2.62	2.60	2.60	2.59	2.59	2.55	2.50	2.46	2.44	2.44	2.41	2.41	2.41	2.43	2.30	7.57	,	
	8.55	0 02	4 11	· •		6	3.68	2.46	0.25	1.18	2.22	1.88	86.0	; ,	3 5	9.00	C/.)	1.36	0.64	36.00	0.75	35 15	0.15	} 6 6	2.65	5 . :	14.62	1.22	1.09	2.61	0.57	1.0	1.50	1.17	2.46	0.94	1.63	0.32	15.73	7.91	0.77	1.38	0.48	21.78	7.4.0		7.77	00.	 	
	30.23	800	3 5	2.0	5.05	2.60	12.10	8.05	0.83	3.79	7.05	5.91	- e	3 3	5 8	9 6	2.32	4.17	1.82	101.29	2.12	08 80		3.6	87.7	48.94	39.72	3.31	2.93	6.96	1.50	2.62	3.92	3.04 24	6.38	2.44	4.21	0.81	39.31	19.49	1.87	3.37	14	52 (2	24.20	ر ان	3.61	3.93	3.10	
	R77948	AA443067	A A 6226 6	A403000	AA41241/	R66101	H09076	R00276	R61372	AA628113	N53447	AA775257	TR7428	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0	7452210 1117076	14/0/6	H0283/	AA418748	R45517	N55459	N59816	A 4 9 6 9 7 9	00000	22,000	H92821	AA487902	T56281	AA878120	T69477	AA496988	N53031	AA150500	T69305	W74533	N74131	H17121	AA702335	R61883	H72722	N34316	AA453526	AA634431	DEARON	704090 LF4340	1000	NZ3139	AA479795	H73590	AA021586	
	445740	04/041	75/200	8565/5	731469	140197	46166	123264	37980	1055707	245277	07050	0/0380	88000	782335	193087	151477	768020	35329	245990	28083	700307	646037	434852	231574	840590	78353	1418621	66902	823655	246430	491751	82905	346583	298417	51178	447416	42872	232772	277.173	705425	74366	43000	9000	202535	267546	740604	214441	364022	
	ç	7 , 9	2 :	.	42	₹	4	45	46	47	. 8	ę s	7	3	2	23	53	2	55	¥	3 6	3	8	20	8	5	62	63	2	65	99	2 2	Š	8 8	8 2	2 2	2 :	: 2	2	, K) q	19	ះ វ	2 1	6/	8	∞	82	8	

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+ 4 + 4 % 0 + 4 % + 8 4 4 4 % + 8 0 % % 4 % % % % % % % % % % % % % % % %	3.15 1.11 0.61 0.49 0.48
2. 2. 3. 3. 3. 3. 3. 3. 3. 3. 3. 3. 3. 3. 3.	3.37 3.19 0.56 0.63 4.04
2.42 2.42 2.43 2.40 2.45 2.40 2.40 2.40 2.40 2.40 2.40 2.40 2.40	0.21 3.72 3.89 3.81
	3.19 3.19 3.18 0.53
8 8 8 8 4 4 8 9 9 8 8 9 9 4 6 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9	0.26 0.26 0.36 0.00
2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2	5.1. 7.1. 7.1. 66. 1.66
0.33 0.34 0.35 0.37 0.37 0.37 0.37 0.37 0.37 0.37 0.38	21.21 27.34 3.24 0.94
0.77 0.84 15.54 15.54 12.27 37.52 34.05 37.52 34.05 37.52 34.05 12.27 13.09 13.09 13.09 13.09 14.12 15.71 15.71 15.71 15.71	37.58 48.33 5.72 1.59 0.03
N67039 AA447522 H08862 T67053 N52151 AA456299 AA456321 AA65331 AA65331 AA65331 AA65331 AA653331 AA653331 AA683345 AA684326 H50086 AA434487 R16596 W49563 W49563 W49563 W49563 W49563 AA609256 AA459292 AA609256 AA459292 AA609256 AA459292 AA609256 AA609256 AA459292 AA609256 AA600224 N80129 AA459292 AA600224 N80129 AA459292 AA600224 N80129 AA459292 AA600224 N80129 AA459292 AA600224 N600234	AA609914
295985 782575 45852 6650 284383 813151 813151 813151 813151 813151 825745 144747 50772 247117 1468260 179083 770868 129585 324815 291097 530814 1032405 49303 1031701 530185 853998 378813 810899 227392 227393	35804 1031113
88 88 88 88 88 88 88 88 88 88 88 88 88	252555

Table 2-9; Good vs. Poor Clinical Outcome: Genes upregulated 5 fold or more in at least 2 tumor samples.

Table 2-9: Good vs Poor Clinical Outcome: Genes upregulated 5 fold or more in at least 2 tumor samples.

		tumor samples.	upies.								1. 1. 1. 1. 1. 1. 1. 1. 1. 1. 1. 1. 1. 1	* Of Timos
	20 4 21	Gen Bank				Fold(AveGood/A	Fold(mpm27/A	Fold(mpm28/A F	old(mpm29/Av	Fold(AveGood/A Fold(mpm27/A Fold(mpm28/A Fold(mpm28/Av Fold(mpm30/Av Fold(mpm30/Av Fold(mpm30/Av Fold(mpm30/Av	old(mpm3a/Av	Samples
Order		Accession	Tissue	AveGood	AvePoor	vePoor)	vePoor)	vePoor)	ePoor}	100.1		
		Number	Distribution				6	4.40	0.27	9.85	7.08	က
-	281039	N47717	bone	38.67	4.42	8.29	75.27	e. -		1 0	08.80	ო
- (00000	0000714	atatata	27 13	3.45	7.87	22.41	0.41	0.67	8.	3 1	. (
7	880282	8079 1 2	prostate	2	4, 0	7.44	8.50	1.97	1.30	9.52	28.6	2
က	155072	R71393	•	3.32	C.4.0	ŧ.;	5	1 98	0.98	5.30	10.17	က
*	208210	H65286	testis	2.70	0.42	6.48	00.4	- 0	6	8.30	5.96	က
10	770066	AA430545	brain	28.72	4.62	5.78	12.05	0.0	- c	990	7.30	က
ဗ	299679	W05628	pone	2.55	0.50	5.07	7.07	6.80	- 77 °C	69	1.02	8
^	767130	AA424535	kidney	4.28	0.37	11.65	1.60	25.69	#*·07		1.22	8
- 60	307337	N95226	spleen	10.81	96.0	11.29	1.65	7.47	6.4.4. Ca. 4.0.	3 40	21.15	8
a	471641	AA034945	brain	11.03	1.08	10.17	21.19	2.20	77.4	2.10	12.92	8
, c	712950	AA282273		38.22	6.9	5.50	8.84	, 2	0.35		7.0	7
: ‡	41889	B6438	adrenal	14.	0.28	5.03	1.46	5.93	15.74	00.1	3 28	8
: \$	84234	T80111		4.13	0.82	5.03	11.18	1.30	0.80	80.00 0	2 5	
<u> </u>		0490514		67 8	0.71	4.90	14.39	0.35	0.46	3.86	76.0	
5	97//97	OCLACN	• .	5		4 56	9.05	2.75	2.93	0.99	7.09	v
*	301735	N90882	esophagus	0r. 4	08.0	6. 6	02.0	3.61	6.33	4.17	6.26	8
5	811162	AA486471	thyroid	28.87	6.85	4.21	2 6	337	0.78	6.56	7.23	7
91	362278	AA001219	lymph	5.53	-	3.91	t 60.71	, q	44.75	0.59	0.31	8
11	1409509	AA868929	bone	2.84	0.75	3.78	0.30	00.0		0.93	0.97	7
8	768953	AA424920		3.43	0.94	3.66	0.96	7.4.C	S. 0	00.6	9.41	8
18	357628	W94121	•	1.80	0.49	3.65	5.62	0.55	9 6	3 54	6.23	8
20	195365	R88904	•	1.81	0.50	3.62	6.03	1,45	0.0 1.0	, K	77.6	М
72	145740	R77948	prostate	30.23	8.55	3.53	1.47	0.52	5 6	. 88.0	1.08	8
23	782335	AA432270	endothelial	* .0	1.30	3.12	6.21	νς 48.	1.80	3		

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatio Epithelial Cell Lines Cell Lines vs. Normal Prostatio Epithelial Cell Lines vs. Normal Prostatio Epithelia Cell Lines vs. Normal Prostatio Epithelial Cell Lines vs. Normal Prostatio Epithelia Cell Li 2 of 3 or 10 fold or higher in at least 1 of 3 cancer cell lines.

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostastic Epithelial Cell Line: Genes upregulated fold or higher in 3 of 3, 5 fold or higher in at least 2 of 3 or 10 fold or higher in at least 1 of 3 cancer cell lines.

Order	IMAGE Clone ID	Gen Bank Accession Number	MPM1(PrEC)	Ave Cancer Cell Lines Values	Fold(AveCell Ilne/MPM1)	Fold(MPM2/M Fo PM1)	.ld(MPM271/M PM1)	Fold(MPM2/M Fold(MPM271/M # of Cancer PM1) PM1) Cell Lines	# of Cancer Cell Lines
	1		5	5.71	591.67	7.33	66.37	1701.31	ო
- :	291/6/	A443850	5 6		319.36	4.57	944.19	9.33	ო
7	510631	AA099431	5, 6	S 99	234 30	8.10	680.66	14.13	က
က	592707	AA160606	5 6	5 6 8	196.56	8.10	570.75	10.82	က
4	220851	H95633	5 6	5 5	180.70	8.88	526.08	7.15	ო
NO.	/82804	AA448180	- 6 6 6	. c	117.17	3.60	344.82	3.10	က
φ	838831	AA481770	0.02	7.00	112 48	13.45	301.13	22.85	ო
^	1343732	AA725564	0.0	7	103.67	60%	302.09	5.84	ო
œ	768226	AA424900	0.0	40.L	95.59	3.74	290.43	3.41	ო
Ø	731048	AA421275	0.01	0.89	n (6		249 63	3.45	က
9	366039	AA071503	0.03	2.66	80.03	- C	239.88	12.54	ო
=	795627	AA459917	0.01	0.51	85.17	5.03 40	203 18	7.41	ო
12	1161830	AA876021	0.01	0.71	71.26	5.18	2000	86.	· e*
13	731371	AA421047	0.03	2.20	71.17	3.28	205.85	4:30) r
4	784296	AA447079	0.01	99.0	65.30	45.84	106.60	45.40	o (
. K.	796569	AA460463	0.01	0.35	63.57	16.10	138.91	33.70	, c
. .	346942	W94289	0.01	0.61	61.20	3.73	168.10	11.76	n (
1 2	705858	A A 4 R 1 5 2 2	0.03	1.60	60.34	6.29	170.43	4.30	m
- 9	131452	072270	0.45	26.93	59.79	104.24	4.69	70.45	ო
<u>•</u>	15 FREEK	A A G7664	800	4.36	54.32	97.06	17.92	47.97	က
<u> </u>	743016	AA406036	0.01	0.50	50.47	15.70	123.80	11.93	က
3 7	30442	R51617	0.10	5.05	49.93	99.52	44.71	5.49	ro (
ī (2307A	D44048	080	36.21	45.35	56.36	29.19	50.51	n
3 8	2000	A 4 9 0 4 0 6	20.0	3.16	45.16	91.62	20.97	22.90	ო
3 3	470003	00180100	5 6	0.44	44.34	25.03	101.07	6.93	ო
7 1	120003	14 / DOC 4	5 6	28.0	43.48	3.54	120.65	6.23	ო
22	1389010	AA833138	7000	0.00	43.37	5.62	121.33	3.16	ო
8	08407	757580	5 5	0.40	41.72	4.59	109.53	11.05	က
27	205/6/	AA437120	5 6	1 0	41.51	3.55	116.08	4.92	ო
78	813633	AA447740	0.0 0.0	1.08	<u>.</u>)) •			

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Lines Genes upregulated 3 fold or higher in at least 1 of 3 cancer cell lines.

367900	AA810225	0.01	0.41	38.18	3.08	99.20 106.34	19.33 5.11	ຕ ຕ (
AA456035	35 5	9.0 10.0	0:30	37.18	16.25	80.79	14.51	ო ი
H17888) 8 8	0.01	0.32	36.79	12.25	87.45	10.68 42.68	n e
H94482	22	0.01	0.36	35.62	5.91	88.28	9.41	, n
N30098	86	0.01	0.25	34.44	20.13 F 81	82.72	11.01	ო
AA608	832	0.0	0.33	33.10	29.44	39.44	28.82	ო
AA045436	436	0.02	0.08	31.52	8.48	9.21	76.87	က
N33331	31	5 6		29 19	24.48	38.30	24.80	က
N72115	2	3 6		29.16	9.10	30.24	48.14	ო
AA863	282	5 6	0.43	29.13	7.73	36.47	43.19	ო
1111	53	5 6		28.83	5.99	65.78	14.72	က
A 783	3328	70.0	1 2	27.34	19.07	21.59	41.29	ო
66H	913	3 6		27.21	13.82	36.04	31.77	m
26Z	764	5 6	2.5	26.71	9.46	65.35	5.30	က
Rec	1981	7.0	- 4	26.24	7.36	35.03	36.33	ო
₹	18903	5 6	2 7	25.96	33.86	30.16	13.87	ო
<u> </u>	818	5 5	1.4.1	25.93	38.89	13.26	25.64	ო
Ž į	234	6 .	22.00	25.84	5.18	58.23	14.11	ო
X :	666	5 6	0.00	25.59	4.23	62.71	9.84	ო
Ž	85136	5 5	0.17	25.59	12.08	52.20	12.48	က
ő š	000	5 6	0.25	25.24	4.46	54.64	16.63	က
	2000	5 6	0.17	25.09	3.61	60.05	11.62	က
3 3	0000	5 6	10.00	24.80	34.18	13.61	· 26.60	က
3	3040 1087E	86.0	9.30	24.51	46.38	16.26	10.88	භ ·
	0000	5 6	0.24	24.29	30.26	10.09	32.50	က
	9000	5 5	0.16	23.97	44.37	17.55	9.98	က
2 3	- 5	5 6	84	23.75	8.09	59.71	3.46	ო
ž :	71.1	5 5	0.00	23.01	8.11	34.50	26.43	က
Ž Š	/60/	5 6		22.37	4.98	30.37	31.76	ო
3 3	M050520	5 6	3.74	22.30	3.82	51.95	11.14	က
Ž	K44830	. c	3.52	22.24	38.20	16.96	11.55	ო
ξÿ	74107	5 6	40.77	21.44	37.52	23.72	3.08	က
8 2	R61866	0.05	0.97	21.43	4.47	56.01	3.81	က
	}			2				

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell-Line Cenes upregulated 3 fold or higher in at least 1 of 3 cancer cell lines.

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47.62	49.72	24.62	22.42	9.30	40.27	43.06	20.16	20.07	- 7.5.	14.13	10.73	16.88	5.21	¥.	4.08	31.84	32.94	19.5	33.48	55.45	04.9 04.00	70.07	40. 40.	12.25	69.9	6.49	19.22	3.61	5.36	15.00	3.17	22.27	5.13	15.54	28 45	26.42	70.44		
10.88	7.89	18 45	33.47	30.67	5 1	0	10.12	24.35	38.83	28.90	33.82	8.91	42.11	11.78	42.48	10.67	8 97	30 yr	90.90	5.09	30.25	15.89	41.27	14.07	36.93	33.79	22.78	39.06	5.66	24.61	37.53	16.46	32.86	20.20	3.02	gr.4	6.51		
4.40	5.07) 0 0	0.10	9.30	20.84	7.04	3.39	10.20	4.49	10.45	7.52	25.95	4.26	, A	4 19	9 10	2 7	40.7	6.72	10.47	9.38	5.81	3.05	21.48	3.94	6.61	4 69	32	33.97	44.4	· 6			6 6 7	24.69	10.07	9.65		
20.97	6.02	20.90	20.42	20.40	20.27	19.50	18.85	18.24	18.01	17.83	17.36	17.25	1 40	11.10	5.5	16.92	68.91	16.56	16.54	16.35	16.34	16.17	15.96	15.94	15.85	7.00 63	0.00	15.57	10.54 0.54	90.5	14.92	14.84	14.61	14.54	14.41	14.24	14.20	m	>
4.0	19.10	0.21	0.25	0.20	2.46	0.19	0.17	0.31	0.35	2 95	2.0	2 40	33.78	0.00	0.63	0.69	0.17	0.17	0.21	0.16	0.09	11.31	1.17		2.0		0.29	0.30	0.71	3.80	0.35	0.70	0.35	0.14	1.64	3.51	0.14		
	0.91	0.01	0.01	0.01	0.12	0.01	0.01	0.02	0.02	0 47	- E		1.96	0.05	o 3	0.0 40.0	0.01	0.01	0.01	0.01	0.01	0.70	200		0.02	0.03	0.02	0.02	0.05	0.25	0.05	0.05	0.02	0.01	0.11	0.25	0 0	;	
	N32904	W90036	H22824	N32542	H22563	R54036	R54492	44883187	AA057105	08170000	AA047441	AA701315	T49355	AA609304	N47012	AA192784	AA676998	W81432	N26769	W92263	U70466	1103717	7 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	H81554	T69270	AA410437	AA418545	AA045175	AA401378	AA702335	H97140	AA406061	R40434	AA437142	AA476543	AA883800	D30064	10000	
			51581																																785293	1461609	75045	01807	
	\$	65	99	2	, «C	8 8	3 8	2 ;	. 1	2	2	4	75	92	#	78	62	80	2	5 6	3 8	3 3	\$:	ဗ္ဗ	8	87	88	88	06	8	85	66	\$	56	g	3 6	6	Ø D	

.25	.35 3	3.16 3	.12 3	.48	3.74	3.62	7.41 3	2,12 3	3,45	.12 3	3.18	.86	.18	26 3		, te); ;	5.06 5.06	S 0	7,55	3.17	3.88	3.11	3.72	3.09	3.40 3	3.89	1.11 3	3.94	.94	3.02	92 3			. F
					•													•								20.61 3.4									
																										10.87									
14.14	14.14	14.13	13.78	13.76	13,71	13.65	13.58	13.43		13.24	13.23	13.22	13.14	13.14	12.88	12.67	12.37	12.30	12.18	12.10	12.06	11.98	11.80	11.73	11.67	11.63	11.61	11.46	11.30	40.87	10.07	10.81	10.70	10.66	
0.50	4 c. 4	17.58), 39), 39	0.00	15.63	16.52	10.52		0.13	8.63	F8.F	0.26	0.23	1.05	0.17	0.25	3.56	0.12	14.39	0.31	16.58	16.82	3.89	034	11.24	0.20	27.64			n a	5.68	90.0	0.09	5.93	
2	5 6			3 6						_	_											·				0.02	•				_	_			
	•		AA774665	K42600	K26149					W93379														0		N64610		•	•		•	•	-	•	
	1031446	32109	856289	30806	41406	08/6501	898286	502003	359269	415089	1466409	824074	51103	1467166	244050	1055121	897807	845441	686172	4040070	785368	764640	40104	10404	1050780	257438	207 130	366971	609228	210820	795746	795840	788209	767403	
	o	5	5	102	5	\$	105	8	107	108	5	110	Ξ	112	113	114	4.5	. .	7 4 4	- ;	2 5	<u> </u>	2 3	<u> </u>	2	2 3	47	125	126	127	128	129	£1	듄	

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line. Genes upregulated 3 fold or higher in at least 1 of 3 cancer cell lines.
2 of 3 or 10 fold or higher in at least 1 of 3 cancer cell lines.

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5.03	4.82	22.97	5.78	14.06	4.93	16.03	15.10	6.87	14.13	18.44	7.37	3.91	5.79	11.50	7.50	9.62	18.79	15.70	8.67	7.35	5.24	4.33	11.30	5.84	4.46	13.62	5.71	13.08	6.90	10.11	11.70	5.53	15.48	9.84	
5.77	22.27	4.07	20.48	3.07	3.47	9.94	8.56	18.90	9.81	6.58	10.28	20.10	10.79	11.42	14.40	13.22	5.97	7.97	15.51	11.24	12.44	10.86	11.63	8.66	14.83	8.45	8.47	10.73	12.47	7.21	9.17	13.78	6.65	10.99	
20.21	3.65	3.39	3.74	12.71	21.35	3.40	5,69	3.45	66.4	3.83	11.04	4.57	11.98	5.53	6.47	5.48	3.48	4.56	3.79	9.36	10.06	12.49	4.58	12.99	8.17	5.38	13.21	3.46	7.38	9.14	5.30	6.69	3.86	5.02	
10.34	10.24	10.15	00.01	995	20.0	9:35	97.8	9:70	9.64 1.00	9.62	9.57	9.53	9.52	9.48	9.46	9.44	9.41	9.41	9.32	9.32	9.24	9.22	9.17	9.16	9.15	9.15	9.13	60.6	8.92	8.82	8.72	8.67	8.66	8.62	ည
7.64	100		0.70		-:04	9.07	0.0	0.52	12.20	. c		31.33	10.61	1.40	2.04	10.53	7.14	25 28 25	0.19	3,43	1.03	0.97	68.6	0.60	3.21	4.	1.40	0.26	0.11	0.05	4.90	12.90	48.92	3.90	
7.0	t 5	5 6	0.0	\$ 5	- 1	rc.0	9 6 6	0.03	07.	7 00	5 6	30.0	5 -		22.0	1 2	37.0	0.62	0.00	0.37	5 6	0.10	80.2	200	0.35	0.16	0.15	0.03	0.01	0.01	0.56	1.49	56.5	0.45	;
1000	AA6/8295	AA626028	AA291556	AA598787	AA625979	R44734	A1003636	AA041197	AA682533	AA157261	AA156342	AA/01652	N332/4	N47.1.13	N30779	NOZOGA S	4 6 2 6 3 4 6	AA304830	W74377	A A 0.4 E E E	A04389	A A 705047	A 446462	70440	110100	H46476	AA598817	R69584	W49672	T50699	AA187351	A 4 1 3 7 0 7 2	HOKA38	AA520999	
	430928	745019	725308	898073	745490	34.134	391987	376452	431242	590500	77/086	433307	010000	676002	303039	250300	700043	45037.5	34665	480243	403213	461488	101101	79764	46604	177884	897956	141726	324901	78217	624627	401184	242062	826273)
•	<u>\$</u>	135	136	137	138	139	40	141	142	143	1 ;	145	ę ;	, , ,	5	n (2 :	<u> </u>	132	2 1	<u> </u>	- + C 4	2 5	137	0 4	60 9	5 5	<u> </u>	. £	3 3	4 A	2 4	2 5	5 5	3

. Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line Cell est 1 of 3 cancer cell lines.

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3.90	14.40	6.83	4.37	8.51	7.71	14.93	15.23	15.35	13.11	4.58	5.12	6.49	10.12	4.46	10.00	11.39	10.84	13.28	11.71	9.05	7.97	11.50	7.15	8.23	5.28	4.93	12.46	12.18	7.97	9.46	14.01	4.48	10.54	
14.21		9.00 3.30	16.37	11 10	13.64	5.10	86.6	2.5	7.53	15.49	6.62	11.97	9.24	7.24	. K	4 17	1.1.7	. r	00.0 00.0	0.00 20.00	60.7	5.10	4.96	10.88	4.41	6.94	4.68	6.84	6.65	6.44	4.60	13.99	7.03	
7.42	4.95	3.80		4. n	7.0	t 46		0.00	4.08 4.08	3.42 3.96	38. t	7 8 .	9 6	11 54	<u> </u>	4.4.5 5.4.5	cc:/	09.4	6.23	5.50 6.80	9.50	7.7	10.53	5.7	12.64	10.35	503	90 F	230	5		3.23 3.75	4 11	; ;
8.51	8.35	8.32	8.31	8.30	8.30	8.13	8.13	8.09	8.05	8.02	8.01	7.87	777	7.77	7.75	7.72	7.70	7.70	7.70	7.68	7.61	7.61	C:07	C:27	56.7	7.44	1 6	7.39	7.54	1.51	06.7	7.28	1.24	9
0.14	0.19	4.39	1.57	0.08	7.83	0.25	21.34	10.44	61.74	0.28	5.63	0.54	2.89	0.60	5.81	0.35	24.59	13.42	1.68	261.61	5.69	5.53	0.15	0.0	5.64	0.20	21.88	23.90	57.27	3.90	0.12	3.80	0.41	5.80
0.02	0.02	0.53	0.19	0.01	9.0	0.03	2.63	1.29	79'2	0.03	0.70	0.07	0.37	0.08	0.75	0.05	3.19	1.74	0.22	34.04	0.75	0.73	0.05	0.0	0.75	0.03	2.95	3.23	7.80	0.53	0.05	0.52	90:0	0.80
T52311	AA485265	AA455786	R60019	W73790	AA425755	W96216	R41329	H54752	T65902	AA705825	H29521	AA455938	H29897	AA625764	AA431741	AA906257	T96718	T55607	W15305	AA676460	R96941	AA608568	N30372	AA070331	AA434403	T61122	AA701455	N93924	AA488367	H29227	AA477400	N70714	H16833	T62072
71977	840150	809557	42906	344134	773383	358599	30148	203275	80399	1292182	52741	813271	190325	745393	782259	1505919	121251	73596	322461	882510	200402	950690	260035	530820	770879	83666	435076	309288	843163	52704	740620	294273	50787	85670
169	2 5	2 5	172	173	174	175	176	171	178	179	180	181	182	183	2	185	186	187	188	189	190	191	192	193	1 2	195	196	197	198	199	200	201	202	203

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line. Genes upregulated 3 fold or higher in at least 2 of 3 or 10 fold or higher in at least 1 of 3 cancer cell lines.

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.,	**	.,	•,	.,	.,	••	.,	•	•	•	**	.,	.,	•	•	• •	•••	••	.,	• •	••	.,	.,	•	ν,	•••	•••	••	••	•	•	•	••	••	
5.38	8.99	8.78	10.69	3.65	7.98	6.50	9.20	11.35	3,14	9.86	3.89	5.69	5.19	12.11	12.23	5.38	6.30	4.64	10.97	7.65	4.14	5.60	6.95	8.44	7.62	8.12	4.80	12.51	3.05	4.63	5.67	11.00	10.01	7.97	
11.90	9.60	7.63	6.35	5.87	7.83	9.95	7.07	6.52	11.26	4.41	3.38	10.31	5.36	4.92	5.03	10.24	4.78	9.34	3.78	4.68	8.01	5.02	9.66	4.39	4.47	3.71	7.44	3.32	5.69	9.69	8.97	4.30	5.16	8.32	
4.39	3.01	5.15	4.40	11.72	5.35	4.65	4.74	3.01	6.40	6.49	13.48	4.76	10.00	3,47	3.20	4.75	9.29	6.37	5.57	7.84	8.02	9.45	3.37	7.13	7.85	8.09	7.52	3.80	10.89	5.27	4.93	4.19	4.26	3.07	
7 22	7.20	7.18	7.15	7.08	7.05	7.03	7.00	96.6	6.93	6.92	6.92	6.92	6.85	6.83	6.82	6.79	6.79	6.79	6.77	6.72	6.72	6.69	99.9	6.65	6.65	6.64	6.59	6.54	6.54	6.53	6.52	6.50	6.48	6.45	^
0000	33.93 75	9.73 18.03	5 6	14.78	3 05	8 6	100 37	0.44	5 0	69 K	27.15	7.46	0.70	2.56	8 14	80.0	26.0	893	27.70	0.30	3.61	2.07	65.62	18.79	4.67	4.52	0.07	35.52	0.42	7.31	2.64	22.29	8.44	7.26	!
9	8. c	1	5 5	. 6	3 6	7 6		2 2	3 6	3 6	9 6) d	5 5	0.37	1 19		50.0	430	4 0 4	9 9	0.54	0.34	. 4 . 6	28.5	0.70	0.68	0.01	5.43	0.06	1.12	0.41	3.43	÷ +	1.13) : :
!	H97488	AA460952	A4665310	AA450011	N45236	7,00344	K/4203	AI015558	VV60180	K36046	N20303	704774	AA405250	NETTO	002066	A A 2 K B 2 3 4	H71230	200000000000000000000000000000000000000	AAAAAAAA	H17139	AA609723	C 2000	A 466603	A 6 600002	AA456655	H59203	AA884622	A4397813	H05645	H20847	NABOZE	111860	H17063	AA045508	201
	251936	796114	80000	/00800	20213	120021	145500	105/345	905014	40773	245575	04 4005	014667	746660	434346	21212	214604	46004	350440	51083	1031885	134710	044746	050445	813149	204214	1468597	725454	43662	51460	281723	E01133	160233	487207	•
	20,	205	902	2	80, 5	502	210	112	212	213	4 4	2 2	917	717	0 7	B C	27 6	- 6	77 6	3 6	22.5	336	27 6	77 6	330	230	3.5	- £	233	33.	3 6	600 000 000 000	73 7	736	3

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatio Epithelial Cell Line: Genes, upregulated 3 fold or higher in at least 1 of 3 cancer cell lines.

ო	က	ຕ	m	· m	•	e e e e e	· •	? (, co	က	က	ო	ო	က	က	ო	(1)	e er	> «		n (n (n (m ·	ന	က	ო	က	က	က	က	က	ო	· en	۰ «) r	3	
9.48	9.25	4.22	4 86	663	3 45) · · ·	P. 4.	8.69	10.52	4.60	4.01	4.07	8.79	90.0	5.79	4.83	4.48	4. 6	9.00	7.86	7.19	5.21	5.37	5.40	5.52	7.53	3.97	9.64	9.01	3.11	7.60	82.6	8 7.4	. 4	0 0 1 4 1 4	86.7	
5.78	707	, e	, d	, d 1	9. i	12.00	3.05	10.41	6.93	5.05	10.49	3.15	10.33	5.59	386) u	0.00	08.0	4.21	5.73	6.77	4.55	6.93	8.09	3.03	7.41	3.15	3.97	4.55	4.93	9.49	5.43	446) ; ;	27.0	4.78	5.30	
80	9 6	9.00	8.52	10.68	6.67	3.73	4.19	3.56	3.26	3.19	3.63	11.55	4 27		n 0	6.00	6.92	7.62	9.63	7.47	3.55	6.43	6.00	4.66	9.62	5.03	7.03	696	3.43	787	4 97	. 4		 	3.50	3.49	4.53	
ç	54.	6.41	6.38	6.37	6.36	6.32	6.31	6.30	6.29	6.25	6.24	22	, c	0.43	6.19 9.19	6.17	6.13	6,12	6.11	6.10	90'9	6.06	6.05	6.04	6.02	, v	5 6 C) @ ; u		. 0 4	9.07 9.86	0 0	5.80	5.83	5.81	5.81	5.80	ω
ì	1.70	0.82	4.82	0.62	47.65	0.47	2.79	90.0	4.53	2.35) Y	2 %	51.70	40.75	12.17	4.70	7.33	0.14	3.47	16.66	4.93	0.08	29.59	7.89	9 6	0.00	5 6	 0 0	1.30 0.40	2 (3 (0.06	2.75	8.00	12.98	43.39	14.21	10.85	
	0.26	0.13	0.76	0.10	7.49	0.07	44.0	6		1 0	o 6	0.02	9.0g	6.55	1.97	0.76	1.20	0.02	0.57	2.73	0.81	0.01	4 89	} •	5. 6	0.0 91.0	5.0	0. 90.	0.24	0.07	0.0	0.37	1.37	2.22	7.47	2.45	1.87	
	AA034115	AA757564	AA431753	W68162	H02231	N50827	PA1972	*******	M444/33	N03/44	R11605	R38640	H84871	AA460685	AA447662	R68581	R10284	H18017	AA405800	A A 425302	L06302	13033£	********	A4458510	AA425404	AA620437	R93875	N29918	N81049	AA453714	H09065	AA099523	W94009	H06113	AA454585	R56870	N59136	
	430007	395955	782476	343298	151055	280907	32050	2020	480797	292936	130004	22895	249603	796694	813586	137890	128947	50983	742115	773073	CIOCII	70000	00000	005418	773147	951068	275871	271076	301104	813827	46154	489594	357531	44255	809535	41198	287637	,
	239	240	241	242	243	244	, ,	Ç ;	246	247	248	249	250	251	252	253	254	255	256	3 1	/67	807	RC7	260	261	262	263	264	265	266	267	268	269	270	271	272	273	ì

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in at least 1 of 3 cancer cell lines.

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6.22	60.8	29	3 5	Q+	t	4.86	4.93	7.68	8.24	77.8	;	- : - :	5.48	10.06	6.45	3.10	5.02	45.4	. A.) i	7.85	7.59	4.14	5.87	7.27	6.98	5.55	6.26	6.39	9 14	06.6	6.6	0.4	3.07	3.08	5.40	7.68		
7.27		0 6	6.03	3.12	3.61	5.66	3.99	5 70	> Ç	0	50.0 50.0	4.25	4.48	3.47	4.88	3.75	883	00 6	76.0	67.9	9.6/	3.50	5.09	5.49	5.58	3.79	5.22	1 6 1 6	7.08 70.4	, c	(3. c	 		4.32	3.79	4.68	5.01	3.47		
3 00	3.95	3.39	4.97	3.75	10.19	6.56	. 8 14	· ·	3.00	4.60	5.49	3.52	6.88	3.30	5.49	6 6	- 89	5. t	8.13 CL :	3.62	3.31	4.98	3.64	6.63	4 66	200	3 6	- (n n n n	D 60.0	3.00 3.00	45.4 45.5	3.21	60.9	6.79	8.25	7.86	5.19	4.37	•	
6	5.80	5.78	5.76	5.76	5.72	26	S 9	0.00	5.66	5.65	5.63	5.63	5.62	1 1	. u	- G	n :	رن 40.0	5.54	5.49	5.45	5.45	5 44		, t	70.0	5.37	5.37	5.34 4	5.33	5.32	5.32	5.30	5.24	5.24	5.21	5.20	5 t 4	<u>:</u> (ဘ
	32.21	13.63	0.16	237.42	3.53) i	5.4	5.22	5.08	15.89	1.94	8.72	26.20	20.50	\$ 10 \$ 0	5.38	2.53	0.40	24.34	5.08	2.80	8.62	5 6	- i	5.65	1.59	11.60	10.19	0.55	0.94	35.80	6.70	13.64	0.10	0.91	. 65 60) t	0. 15 7. 15	83.75	
	5.55	2.36	0.03	41.24	0.83	0.02	0.83	0.92	0.90	2.81	0.34	1 A A	3 9	94.4	1.43	96.0	0.45	0.07	4.40	0.93	0.51	. 4 8	00:-	0.00	 4	0.30	2.16	1.90	0.10	0.18	6.73	1.26	2.57	0.02	0.17	- 0	0.03	0.02	16.19	
	AA625662	T57359	H04332	A A 6 6 0 3 4 1		AA460310	AA150777	R81486	AA228130	AA629849	H08785	2010011	AAZ/8990	R80790	AA733061	H93424	R25614	H23983	171965	AA115559	N120503	2602N	K42685	AA017213	AA598803	AA779383	AA450205	AA682545	AA676840	W93682	AA620556	N49079	H98822	BOOK28	041760	K6//36	AA019511	R52085	AA485898	
	745360	74738	447385	00741																									460114				261714	26207	16707	18081	363103	40299	840506	•
	27.4	1 1	0.75	0/7	//7	278	279	280	281	, ac	3 6	287	28	285	286	287	288	289	200	200	187	292	293	294	295	296	297	298	299	300	301	302	203	3 6	3	302	306	307	308	

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes Upregulated 3 fold or higher in at least 1 of 3 cancer cell lines.

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6.97	5.45	5.93	4.75	8.15	4.97	4.67	5.63	4 45	S:+ *	, 4 , 4	2 2	5 4 5 5 4	9 4	04. A	A 25	2.00 2.00 2.00 2.00 3.00 3.00 3.00 3.00	0.0	- 6 6	08.c	4.80	5.64	6.54	4.37	6.23	4.69	3.60	5.29	3.08	3.05	3.31	6.55	5.01	3.64	, i	5.5	4. 59.	
5.27	5.00	5.12	4.23	3.76	3.23	7.05	3 6	50.0	9.00	o.4.	0.4	0.70	0.0	6. i		4.74	5.83	3.84 4	5.34	4.57	5.49	4.24	4.27	4.29	3.94	3.35	4.33	7.24	4.15	4.89	3.95		2 2	9.53	7.07	4. 46.	
3.25	4,99	4.34	30	9 Y	i i	7.10	3.02	4.62	4.64	6.45	4.26	4.49	3.03	6 .00	5.36	3.28	3.32	4.46	3.38	5.12	3.32	3.64	5.78	3.85	5.62	7.29	4.52	3.82	6.91	5.9	2.56) ¥	2 6	5.03	3.32	4.03	
5.16	5. 15.	, r	5 4	5.12	5.12	5.12	5.11	5.09	5.06	5.06	5.06	5.02	5.02	5.01	4.96	4.95	4.92	4.90	4.87	4.83	4.81	4.81	4.81	4.79	4.75	4.75	4.71	4.71	0,4	07.4) (i	D (4.07	4.65	4.63	4.63	10
2 47		<u>.</u> 6	31.82	0.40	0.24	34.01	2.84	0.77	10.81	23.18	8.96	1.02	23.47	4.55	0.75	24.26	11.53	6.70	2.66	1.54	0.90	0.81	15.65	1.22	4 83	7.74	. v) ; 6	5 6	32.03	5.63	0. J	3.48	1.69	3.36	0.43	
87.0	£ 6	10.	6.20	0.08	0.05	6.64	0.56	0.15	2.14	4.58	1.77	0.20	4.68	0.91	0.15	4.90	2.34	1.37	0.54	0.32	0.19	0.17	3.26	3 6	55.5		07:-		t 0	6.94	0.78	0.0 4	0.74	0.36	0.73	0.09	
	AA136060	AA703000	N66132	R80387	AA663995	AA450265	H18068	AA885851	AA463924	N93715	AA448160	AA683557	AA504272	AA129552	AA626784	N40939	AA443969	AA459632	AA418918	A A B 78135	AA156599	A A 776602	A A 620662	A4620333	AA481/5/	STORN	N50745	AA88312/	AA621019	AA448396	N73301	H97000	T66936	AA046321	H23482	AA682405	
	490060	447208	278483	147147	855390	789182	50649	1500388	810264	307119	782766	505864	825411	564803	877767	277163	757206	810974	767994	121805	5000 E	4202020	054443	201142	68/018	300140	283/51	1467026	1029501	781341	292068	249949	66406	376764	52543	461775	
,	308	310	311	312	313	314	315	316	317	348	349	320	321	322	323	324	325	328	32.4	770	0 00	876	3	555	332	333	934 4	335	336	337	338	339	340	341	32	5 2	ŝ

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Lines Genes, upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

ო	က	က	က	(1)	er.	۰ «	· •	n (ro (m	က	က	က	r)	က	က	ĸ	က	ო	က	က	က	67	· "	, c	, (n (n	က	က	က	က	က	ო	က	e-7)	
7.60	5.45	5.80	5.89	8	4 05	4 6	6/:0	6.37	5.59	5.44	5.34	4.18	7.01	4.75	4.30	3.12	3.19	3.76	4.54	4.00	3.87	5.27	83	70.0	07.4	ο ·	5.10	4.99	3.59	3.19	5.47	4.18	5.33	3.24	5.43	2,60	ĝ.	
3.22	5.08	3.14	2 K) () ()) h	0.70	4.81	3.27	3.51	4.39	5.08	6.13	3.41	3.36	5.80	5.13	4.50	5.42	5.36	44.4	3.81	9. 4		00.00 00.00 00.000	4.5	3.76	3.33	5.05	4.31	6.72	4.21	4.72	3.51	6.39	724	, C	R.	
3.06	3.28	4 85	, 4 20 C	4. c	6.38	3.02	5.14	4.05	4.56	3.79	3.16	3.27	3.14	5.43	3.35	5.20	5.70	4.16	3.43	4.85	, A		2.05	3.03	5.77	3.23	4.72	3.06	5.17	3.11	3.32	4.06	4.10	3.30	767	t 7:	3.40	
. 4.63	4 50	 	0	9.40	4.58	4.57	4.56	4.56	4.55	4.54	4.53	4.52	4.52	4.51	4.48	4.48	4.46	4.45	4.45	0 F F	t - 4 5 - 6	24.4 24.4	04.4	4.39	4.39	4.38	4.38	4.37	4.36	4.34	4.33	4.32	4.31	4.34	- G	05.4	4.29	Ŧ.
22 62	44.07	70.11	3.08	8.14	0.99	68.72	0.0 40.0	25.23	5.67	16.81	13.59	19.58	0.60	3.85	133.80	10.51	5.43	40.45	20.43	32.01	2.30	2.19	10.04	4.57	7.88	2.06	0.93	0.18	9.99	0.09	17.91	. 79	5.41	. 4	7.03	5.44	1.66	
6	B 6	2.40	0.67	1.77	0.22	15.03	0.01	5.53	1.25	3.70	3.00	4.33	13	38.0	20.02	2.35	, t	77: 0	6.35	ર ક	0.67	0.40	2.28	. 2	1.80	0.47	0.21	0.04	2.29	0.02	4.13	, C	 	24:-	0.62	1.26	0.39	
1	AAU/4666	AA173926	N69491	N62245	AA699762	N74956	H43317	AA48184	WOORS	MEDOZO	A 4 5 7 8 7	AA056305	A 4 70 7 70 4	25,300	A A 68208F	A0000000	A4454347	AA126/39	AA453273	W96114	W84486	AI014704	W60015	AA486430	AA857804	AA010128	R94633	H84926	N95495	AA284108	N140284	1049401	60000	08080	W90164	N30185	R12804	
	544564	595109	292933	287749	461235	295551	186918	782800	206155	202200	588840	500040	200413	10014	397.90	90000	0.4700	940049	785371	358457	356732	1627623	341942	842906	1475463	359135	198104	249562	308466	324313	201212	4004	1000+	81 COC7	418150	258761	25274	
,	¥	345	346	347	348	349	350	35.4	26.	400	200	# H	0 0	6 i	<u>ک</u> دو	000	PG:	360	361	362	363	364	365	366	367	368	369	370	5 5	3 6	7 6	2 2	4/5	3/3	376	377	378	

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelia! Cell Line Genes upregulated 3 fold or higher in at least 1 of 3 cancer cell lines.

က	က	က	က	က	က	ო	က	က	က	ო	က	ო	ო	ო	က	ო	က	ო	က	က	ო	က	က	က	ო	က	က	က	ო	ന	ო	က	က	ო	•	
4.31	5.24	4.93	5.79	3.61	4.55	3.43	3.75	4.67	3.12	4.67	4.40	5.97	3.26	5.22	5.18	3.14	5.01	4.22	3.31	3.26	4.71	4.63	5.40	3.23	4.30	5.02	5.93	3.07	5.07	3.44	4.63	3.86	5.33	3.59	}	
4.70	4.54	6 6 6	3.05	3.07	5.07	5.59	5.64	4.51	5.97	4.63	4.75	3.59	. 60 60 60	3.76	3.72	4.62	04	, 4 9 80 80	50.4		- 60 60 60 60 60 60 60 60 60 60 60 60 60 6	0 0 0	85 85 85 85 85 85 85 85 85 85 85 85 85 8	5.27	4 59	4.10	3.13	3.14	3.29	474	3.60	7.7.	 	, c	0.6	
3.87	3.03	9 6) ()	06.0	90.0		3.06	3.45) (40.0 08.6	3.43	9 6) (r))) (S 6		4 c	ن د د د د د د د د د د د د د د د د د د د	6.10	4	9. 4. 4.	t 99 c	00.5 80.00 80 80.00 80 80.00 80 80 80 80 80 80 80 80 80 80 80 80 8	5.50	7 .0	6. c	3 6	, r	 	4 6	. c.	21.0	7.5. 7.4.6	رن 10. دو	3.30	
7 30	57.4	4.27	4.25	4.25	4.23	4.22	4.22	4.22	4.21	02.4	0.4.4	4; 4 D (n (81.4	0 4	4.16	4.14	4.13	4.13	4.12	. 1 .	1.4	90.4	80.4	4.07	4.07	, 4, 4 , 70, 4	50.4	4 4	4.03	4.00	99.50 19.50	3.97	3.97	3.96	12
i i	2.72	8.22	0.82	1.80	7.01	1.26	9.68	9.83	7.07	15.01	1.52	0.45	5.38	9.46	5.68	0.22	6.85	26.51	1.14	5.90	21.16	2.02	30.82	2.60	72.78	4.98	8.49	8.55	10.05	3.22	0.34	9.11	16.49	32.97	15.20	
	0.63	1.93	0.19	0.42	1.66	0.30	2.36	2.33	1.68	3.57	0.36	0.11	1.29	2.26	1.36	0.05	1.66	6.42	0.28	1.43	5.14	0.49	7.53	0.64	17.87	1.22	5.09	2.11	2.49	0.80	90.0	2.29	4.15	8.31	3.83	
	AA251826	W88792	AA975388	AA126259	AA679569	R59722	H44448	AA024637	W47254	AA459909	N92478	AA489247	AA025937	T59668	AA452966	AA496360	AA838691	AA053129	R08935	AA644099	N89861	AA125825	R05693	N64617	AA678021	AA489000	R33303	H23985	AA705142	N64508	AA460571	W86860	H01926	H11792	N29901	
	684835	417855	1556056	502767	453599	42793	183771	365157	324255	809421	301867	825089	365641	76196	788566	755301	1387760	510170	127925	845380	305538	548693	125183	289945	431803	824739	136180	51469	461592	290378	795830	416390	150135	47681	271006	
	379	380	381	382	383	384	385	386	387	388	389	390	391	392	393	394	395	396	397	398	388	400	401	402	403	404	405	406	407	408	409	410	-	412	413	· ·

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithellal Cell Lines Genes, upregulated 3 fold or higher in at least 1 of 3 cancer cell lines.

			ŏ	3.60	3.96	4.14	4.14	3.60	` ຕ
414	625/64	AA188366			3 94	3.65	3.46	4.71	જ
415	1654974	A1023724	2. 40.	9.05		4 53	3.70	3.55	က
416	194305	H50656	0.29	1.14	3.80 0.00) ·	4.45	4.15	က
417	32870	R43205	0.34	1.32	3.93	9 9	2 5	3.57	က
418	786083	AA448676	2.27	8.89	3.93	4.14	5.6	3.47	· 67
2 4	613126	AA211448	0.43	1.69	3.92	4.47	3.83	ř	. «
1	202676	1160203	4.31	16.90	3.92	3.28	3.75	4.14	· ·
27	0.0007	T05607	0 02	60:0	3.86	4.33	3.12	4.12	n (
421	007011	120001	20. E	11.74	3.85	5.38	3.09	3.09	n (
422	754355	AA436152	9 6	40.26	3.85	4.86	3.02	3.67	က
423	77244	T50139	9 6	10.20	48.6	4.81	3.36	3.36	က
424	813675	AA453832	2.30	2 !		4.55	3.33	3.58	က
425	295710	N72697	1.17	4.47	5.05 5.05	3.23	4.39	3.59	ო
426	884690	AA630016	0.01	0.0	\$ 100 d	3 24	3.48	4.65	m
427	377884	AA777092	0.25	0.95	3.74		4 03	3.19	es
428	453642	AA775888	0.09	0.3 4	3.68	5.65	. 4	3.51	က
429	1586340	AA974348	2.85	10.51	3.68	4.08) i	98	e
430	757404	AA426341	5.69	20.94	3.68	3.33	6.00 6.00) .	er.
7	745286	AA625558	0.16	0.59	3.66	3.24	3.30	? c	, r
·	772500	A 4 2 0 4 2 2	2.41	8.80	3.66	3.28	4.40	3.28	, (
432	BBCC/	77467474 77478474	. L	5.73	3.66	3.48	4.23	3.25	m
433	48/1/23	AA04335		0.05	3.65	4.71	3.21	3.02	က
434	288658	N79353	07.0		3.64	4.05	3.12	3.76	က
435	279920	N57526	60.0	15.3	3 64	3.48	4.25	3.19	က
436	813648	AA453679	3.91	14.21	. c. c	3 22	3.26	4.37	က
437	136744	R35230	5.00	7.22	3.02	27.6	3.44	3.70	က
438	487425	AA046523	2.71	10.00	5.0) (i	3.72	3.07	က
439	589479	AA157129	4.22	15.14	3.58	5.50 5.00 5.00 5.00 5.00 5.00 5.00 5.00	3 65 K	3.62	ო
4	256260	H94617	4.40	4.95	3.54	50.5	5 c	3.24	က
4	627273	AA191461	1.01	3.54	3.52	80.0) i	3.32	n
442	1604596		0.80	2.80	3.49	3.29	\$ G	# G	· «
773	843224		5.65	19.64	3.47	3.47	3.36	6.09	•
? :	784230	AA48394	0.19	0.64	3.42	3.04	4.21	3.02	? (
1		200320	22.0	2.45	3.41	3.42	3.22	3.60	m
4 0	27,000	SDC SE	; u	10.81	3.41	3.01	3.43	3.80	m
446	840384	AA485773	- 60	3.63	339	3.46	3.12	3.59	60
44	412949	AA707840)O'L	3.02	3.37	3.82	3.16	3.13	က
448	731044	AA421273	1.36	70.4	5.5	<u>;</u>			
					13				

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostate Epithelial Cell Line Genes upregulated 3 fold or higher in at least 1 of 3 cancer cell lines.

က	က	7	7	73	7	7	8	2	. 6	2	. 6	. 61	8	. 7	7	8	. ~	, ,	4 (NE (7	64	7	7	7	7	8	7	7	7	7	7	7	7	74	1	
3.05	3.17	7.96	7.05	1917.92	5.67	6.10	2.42	4 72	1 to	5.7.7		33	. 6		146.14	105.59	13.76	13.70	C.4.	99.88	0.73	123.70	0.72	1.93	5.39	3.40	6.47	2.54	12.17	91.51	1.00	3.10	7.45	4.06	28.6	Ş	
3.55	3.07	2966.06	2506.90	61 14	1831 96	1516.35	1331 80	1331.03	1284.73	56.1711	841.07	67.4.00	407.71	457.34 228.30	336.20	4 0.40	150.45	229.57	205.80	95.72	173.62	51.92	88.50	154.39	150.40	149.73	1.37	145.07	139.36	56.58	142 92	2.7.0	428 30	20.20		123.63	
3.08	3.14	86	3 3		7.7	1.50	14.2	3.24	2.48	0.73	9.	10.58	1.61	3.95	1.4.1	0.80	0.89	1.42	2.42	0.87	6.88	1.03	78.84	5.36	45.0	1 05	144 61	4.68	69.0	8 6	- a	27.00	0.75	47.7	7.97	4. 48.	
3.23	3 13	2 6	991.80	838.64	660.46	613.07	508.28	445.85	433.98	393.88	283.26	228.86	157.55	154.10	114.55	95.78	85.65	81.58	70.86	65.49	60.41	58.88	56 02	53.90	50.50	51.30	50.50 C8 C8	50.02	5.50	20.72	10.04	49.10	47.88	45.99	45.56	44.45	4
y O) (i	0.66	9.92	8.39	9.14	9.00	5.08	18.56	4.34	3.94	2.83	1.70	1.58	1.54	2.74	1.93	6.83	0.82	0.71	5.16	0.83	0.50		14:71	0.70	17.7	3.93	60.94 49.94	χ. Σ	0.10	13.02	0.49	12.64	0.60	0.46	0.87	
4	C1.7	0.21	0.01	0.01	0.01	0.01	0.01	0.0 40.0	0.0	0.01	0.01	0.01	0.01	0.01	0.02	0.02	0.08	0.01	0.01	. 60	8 6	0.0	9.0	0.22	. 0.01	0.0	0.08	1.20	0.19	0.02	0.26	0.01	0.26	0.01	0.01	0.02	
	AA127861	A1024401	N74995	AA009615	R60995	N57483	AA433885	T97352	AA479498	N64379	AA429904	AA46446	AA446866	AA812676	AA169154	H56453	AA485214	W80611	AA476833	AA 170033	AA419460	AA757754	AA251784	W73144	AA086475	R40231	AA873635	AA437224	AA443936	R44741	AA504844	AA453802	AA905896	AA449120	R59936	AA292429	
	501731	1640931	299559	365551	42627	277327	773640	121863	754291	290213	781157	781036	784216	1377071	609935	203878	815794	347182	644273	5/5/10	746152	395708	684879	344589	562811	30207	1358266	757435	757143	33827	825740	813748	1505360	785849	43101	714498	
	449	450	451	452	453	40.4	455	456	457	458	459	460	461	462	463	464	465	204	Ş (467	468	469	410	471	472	473	474	475	476	477	478	479	480	481	482	483))

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelia Cell Line Genes apregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

127.57	123.35	105.49	107.04	103.70	103.43	100.95	52.03	96.63	93.52	89.38	0.12	81.20	34.54	7.97	76.38	9.84	70.25	69.04	1.36	27.83	72.00	11.75	66.04	68.60	67.88	71.30	64.40		41.17	41.17	41.17 26.82	41.17 26.82 60.48	41.17 26.82 60.48 9.54	41.17 26.82 60.48 9.54 28.65	2.32 41.17 31.35 2 2.27 26.82 45.50 2 1.76 60.48 11.25 2 0.44 9.54 62.76 2 0.95 28.65 43.13 2 0.87 21.91 48.25 2
							•																												24.86 1 24.50 4 24.25 23.68
																																		_	0.01 0.24 1.09 26.54 0.31 7.41 0.15 3.51
																																			H96647 T57803 AA490945 AA504625
484 811121	485 743362	•		•						493 //3003		495 49530				•									-			•							

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Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line Genes upregulated 3 fold or higher in at least 1 of 3 cancer cell lines.

7	7	~	7	7	7	7	7	7	7	7	7	7	7	7	7	8	8	8	8	~ ~	· 74	8	8	7	. 8	. 6	8	2	8	8	8			, ,	v		
25.34	0.73	53.51	54.20	23.77	3.87	3.43	17.48	3.78	28.27	20.17	50.17	45.41	25.26	49.74	46.16	1.08	5.54	6.22	80.00 40.00	16.22	8.77	1.27	12.21	24 18	7.22	5.65	28.36	25.15	3.27	0.57	15.13	78 37	5.0	3.12	9:9/		
42.67	61.72	9.53	10.05	1.50	58.79	57.89	42.76	0.20	30.54	38.72	7.58	10.31	30.81	1.00	1.93	14.31	47.06	4 4 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5	45.91	40.44	04.13 40.64		37.48	£ 5	1.0.	39.78	18.52	21.02	41.42	24.14	41.20	20.43	01.8Z	38.63	0.27		
.00	3.92	3.5	88	30 S	22.00	2.0		6.33 7.83	20.50	64:1	5 +	2.79	. c	, i	9 0	9.03	ī (2.02	1.2.1	1.8/	1.97	75.7	46.76	1.8/	14.65	2.86		20.7 20.00 2	9 4	90.C	3.92	56.	0.42	2.32	36.64		
23.00	22 42	1 7 7	24.74	17:17	07:17	21.13	20.73	20.39	20.24	20.04	19.87	19.01	18.30		16.91	19.26	18.93	18.21	17.78	17.62	17.44	17.25	17.22	17.19	16.78	16.62	16.47	15.84	15.64	15.42	15.23	15.03	14.95	14.69	14.60	16	
200	0.50	0.30	4.0	3.65	0.14	0.51	0.95	6.19	1.02	3.94	0.20	3.26	0.20	3.30	0.19	0.19	21.01	0.74	3.45	0.54	0.17	0. 44	8.76	1.06	0.11	0.29	1.14	2.94	5.98	2.60	0.95	3.91	3.94	1.38	0.54		
Ş	0.0	0.0	0.02	0.17	0.0	0.02	0.05	0.30	0.05	0.20	0.01	0.17	0.0	0.17		0.0	. .	9. 4	0.19	0.03	0.01	0.03	0.51	90:0	0.01	0.02	0.07	0.19	0.38	0.17	90.0	0.26	0.26	0.09	9.0		
	170198	H88143	AA161097	AA491261	H10072	W95757	N24848	AA255954	R43910	R95684	AA461508	AA520979	R86970	H28091	AA496022	AA456628	T65736	AA620747	AA487505	R60020	AA001376	AA489662	AA857809	AA857163	AA481481	N26559	N51297	AA504838	AA417713	AA206311	N73083	AA463445	H18932	R49592	AA035745		
	81050	220655	591055	824270	46843	358316	267085	686552	32962	199337	795831	826256	197265	162310	759163	809583	80338	1049284	839048	42907	361899	823727	1475476	1410444	756629	266366	283070	825726	746232	647679	291947	811766	51599	37205	364865	,	•
	519	520	521	522	523	524	525	526	527	528	529	530	531	532	533	534	535	536	537	538	539	540	541	542	543	\$ 4	545	546	547	548	549	550	15.	553	553	<i>;</i> <i>;</i>	

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatio Epithelial Cell Line Cenes upregulated 3 fold or higher in at least 1 of 3 cancer cell lines.

8 6			. 6	8	- 71	7	2	7	8	7	74	8	7	8	8	8	. 7	8		4 6		, ,	4 (7	8	7	8	7	8	7	7	7	8	. ~	•	
2.07	2.53	2 7 6	, 4 , 4	50 6	5.10	27.66	12.18	18.27	29.19	26.50	7.37	25.70	28.39	2.23	5.10	4.25	58	70	t):	20.39	20.30 20.30	6. 7	90.1	4.42	20.10	3.37	16.24	28.00	3.44	14.01	28.79	3.60	3.82	3.43	P S	
38.09		36.42	37.89	33.90	26.36	30.50	76.44	21.58	8 82	. 4. 5. 4. 5. 5. 5. 5. 5. 5. 5. 5. 5. 5. 5. 5. 5.	- C	12.28	8 47	20.52	31.40	4.10	21.13	14:33	29.00	0.39	91.41	23.87	29.43	2.25	14.95	30.06	17.13	4.46	30.95	20.54	0.97	28.84	7 6	t	79.90	
3.43	9.	3.59	1.58	1.89	36.41	8 6 6 6	66.7 6	2.42	2 2 3 3	2.00	20.00	30.00 F	0.00	9 7	\$ \$	 	2.47	22.5/	1.18	10.40	1.96	2.11	6.29	29.85	1.00	2.42	2.08	2.89	0.81	, A	5 7	- 6 - 7	1.92	2.83	1.36	
14.53	14.40	14.24	14.20	14.13	14.13	14.09	13.89	13.68	13.50	13.36	13.00	12.97	12.88	12.81	12.79	12.64	12.63	12.56	12.41	12.39	12.38	12.36	12.24	12.17	12.02	11.95	11.82	11.78	7.1.0	1.7	11.7	11.47	11.45	11.23	11.22	17
5.51	0.14	0.42	2.42	0.77	12.46	2.19	4.17	1.3	3.95	8.41	0.08	0.36	0.13	8.32	0.71	1.99	0.33	0.13	4.99	9.44	0.12	3.72	0.12	18.67	1 69	S :- C	73.62	23.33	21.0	0.12	0.18	0.12	0.70	7.46	2.42	
0.38	0.01	0.03	0.17	0.05	0.88	0.16	0:30	0.10	0.29	0.63	0.01	0.03	0.01	0.65	90:0	0.16	0.03	0.0	0.40	0.76	0.01	0:30	0.01		3 7	÷ ;	5 5 6	33. ·	0.01	0.01	0.02	0.01	90.0	99.0	0.22	
AA863125	AA434388	AA453260	R40105	AA010224	N53380	AA699601	W90705	R37738	AA427719	H15549	N52193	AA035730	AA017379	AA911236	R43915	AA070435	H11051	AA453616	AA437212	AA620697	AA022496	H10713	767077	128101	A440401/	R92201	AA398321	AA477227	W51795	AA456143	AA161188	W72881	H18015	AA478952	AA609300	
1455603	770860	795347																															50982	753984	1031561	
	4 H	66 4	55.7	558	559	280	561	562	563	564	565	566	567	568	569	570	57.5	573	1 (27.5	7 7	37.3	0/6	2/6	578	579	280	581	285	583	584	585	586	587	588	

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatio Epithelial Cell Line Genes upregulated 3 fold or higher in at least 1 of 3 cancer cell lines.

0 0	8		. 6	8	8	8	. 6	۰ ۸	۰ ۵	۱ ۸	. 04	8	8	8	8	8	8	7	8	8	7	8	8	8	61	7	7	8	7	7	8	8	. 8		
16.34	75.76	11.15	13.80	1 0 4	4 4 5 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4	18.62	10:02	3. F.	30.40	20.10 30.00	6.31	21.77	4.40	4.66	2.78	6.78	4:17	15.60	3.50	23.76	15.64	0.48	1.65	1.83	17.42	21.42	1.12	16.30	1.00	15.31	24.03	40.88		<u>.</u>	
16.85	22.78	4. d	1.00	08.71	77.07	75.57	15.37	Z0.Z	27.39	10.22	1.00	000 8	25. 25. 25. 28.	1.37	23.39	21.63	24.49	07.77	5.4.4.0 5.4.6.0	631	14.23	7.24	17.15	25.02	11.85	90.9	20.18	12.15	24.20	13.69	0 0 0	0.30 1.30	70.71	19.71	
0.33	2.56	1.10	3.77	0.92	1.90	2.21	0.51	5.09	1.79	2.05	22.91	2.40	4 ,	41.1	23.10	4. c	/o.2	2.40	0.88	78.7	1 6). C	6.4.0 1. 6.1	1.5	? * 0	2.67	. a	9 4	÷ ;	0.4 0.5	6.6	4.65	1.10	8.07	
11.17	11.06	10.94	10.91	10.90	10.89	10.87	10.84	10.83	10.82	10.79	10.76	10.65	10.50	10.47	10.38	10.36	10.36	10.35	10.32	10.32	10.31	10.21	10.19	רו.טר	10.10	10.07	10.03	/ 6. ci	ය න	06.6	9.89. 6	68.6	9.87	9.86	18
0.34	0.95	1.32	0.11	3.90	0.36	1.12	6.12	0.31	3.13	9.32	0.11	7.96	8.09	7.51	18.95	12.67	0.44	0.21	0.87	4.59	4.21	5.52	9.47	0.12	8.21	2.53	16.05	2.42	2.29	0.10	4.94	45.45	3.82	0.09	
0.03	0.09	0.12	0.0	0.36	0.03	0.10	0.57	0.03	0.29	0.86	0.01	0.75	0.77	0.72	1.83	1.22	0.0 40.0	0.02	0.08	0.45	0.41	0.54	0.93	0.0	0.81	0.25	1.60	0.24	0.23	0.01	0.50	4.60	0.39	0.01	
AA160670	AA427737	A A 454222	N31808	H48070	AA453787	R66438	AA410434	N74131	N66900	N30751	AA680136	R39765	AA489073	AA463248	AA702803	AA421171	N93721	AA418852	AA281784	AA099148	H54451	N49389	AA412049	AA158035	AA775223	R94504	AA478036	AA626167	AA664007	AA430042	AA292226	R25377	N21633	R55220	
592728	770854	705528	271483	193333	813738	41869	753428	208417	287190	257955	433155	26164	824802	797055	448036	731095	307138	768043	712401	489509	203184	277621	729953	590640	868838	197651	740457	745248	855391	781110	725877	133136	266135	154720	
Q	600		- C	1 0	8 8	404	2 4	2 6	a c	0 0	809	60	602	603	604	605	909	607	808	609	610	611	612	613	614	615	616	617	618	619	620	621	522	1 6	770

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostate Epithellal Cell Lines Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

8	7	7	7	8	7	7	8	73	7	73	7	7	7	7	~	7	7	7	7	7	7	64	7	7	2	7	7	8	7	7	7	7	61	7	
0.46	4.63	3.25	5.79	18.56	4.67	6.28	3.39	9.58	10.89	0.37	1.89	15.44	16.97	18.46	21.00	3.86	18.37	1.83	15.06	14.44	21.78	23.74	9.79	17.71	5.45	13.51	14.35	21.12	14.67	5.38	9.46	1.10	4.17	16.60	
23.34	22.72	24.64	22.17	9.16	23.22	21.91	24.53	17.50	15.63	3.74	3.03	11.85	1.00	<u>.</u> 2	5.32	23.58	1.12	21.39	4 2.7	12.53	1.86	1.00	16.72	7.70	19.73	13.04	0.36	1.00	10.87	20.24	15.22	13.45	21.38	0.30	
5.59	20.	1.50	1.35	1.31	1.13	0.66	0.81	1.61	2.05	24.42	23.58	1,15	10.45	8.70	1.84	0.66	8.59	4.86	11.43	0.97	4.26	3.14	1.35	2.30	2.37	0.81	12.50	5.08	1.65	1.54	2.44	12.54	1.42	10.05	
08 6	08.0	97.6	27.6	89.6	29.6	9.62	9.57	9.56	9.53	9.51	9.50	9.48	9.47	9.40	9.39	9.37	9.36	9.36	9.34	9.31	9.30	9.29	9.29	9.26	9.18	9.12	9.07	9.07	9.06	9.05	9.04	9.03	8.99	8.98	19
2	75.0	2.73	0.21	79.5	0.10	6.12	4.60	0.16	0.77	6.70	12.98	1.40	60.0	60.0	5.02	3.84	0.08	0.32	90.0	0.22	0.09	0.09	6.35	4.10	0.87	2.44	0.26	0.09	3.21	3.62	10.62	11.71	4.35	17.81	
6	70.0	t 8	200	3 6	5 6	90	0.48	0.02	0.08	0.70	1.37	0.15	0.0	0.01	0.53	0.41	0.01	0.03	0.0	0.02	0.01	0.0	0.68	0.44	0.10	0.27	0.03	0.0	0.35	0.40	1.17	1.30	0.48	1.98	
4	AA464613	AA262080 546426	K49120	A 620620	A 4 4 7 0 8 6	AA133500	AA487432	R89490	R46000	AA018980	AA032221	AA219060	W16659	AA679180	AA136664	H11519	AA699926	N54254	AA284072	N50962	H57060	AA045074	AA455242	AI017607	R95962	R96579	W44766	AA100696	N26515	AA192527	AA488175	AA172056	AA158346	AA620485	
2000	812884	205001	20080	490204	782728	K12116	841323	105695	35804	362926	375682	630013	320495	866702	490959	48226	435330	247628	700792	281162	204596	487165	814791	1627705	230882	199577	320865	490556	266259	627541	877632	594758	591095	951241	,
ć	624	625	979	/70	070	870 9	950	- 63	433	5 5	635	636	637	638	639	840	\$ 54	CPS	643	94	645	846	647	648	649	650	651	652	653	654	655	656	657	658	

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Lines Genes upregulated 3 fold or higher in at least 1 of 3 cancer cell lines.

7	7	~	7	8	2	7	7	8	8	8	7	8	7	81	7	7	8	8	, ,		1 (4		۱ ۵	, 0	1 0	8	8	8	7	•		, ,	4 (7 (7	
5.40	11.94	8.45	14.43	8.30	18.10	0.61	3.16	7.50	3.36	14.35	18.86	17.99	19.50	11.27	12.36	17.21	2.28	16.24	10.27	13:51	23.00	44.26	25.0	18.07	0.01 89 K	9.48	4.38	17.47	5.36	13.48		9.0	04.0 0.0	Ö.00	3.15	
20.84	2.96	17.16	2 08	± 24	. W	2.52	24.74	- 1.1.7	71.85	8:17	00	 	3.76	12.46	14 84	A. A.	5 th	10:07	0.27	0.41	13.18	1.0/	41.11	19.58	0:36	13.54	7 28	8 .	10 34 20 34	\$.00 .00	80 .	99.7	14.68	13.94	18.08	
0.65	11.72	600	0.92	76.6 96.0	0.30	\$ G	n (0.	0 !	1.87	0.0 0.0 0.0	10.33	77.0	0 9	06.4	2 6	5	1.61	4.16	2.41	5.22	8.74	.03 00 00 00	2.23	4.39	5.83	0.1	7 9	n c	2.0	0.38	2.25	0.81	0.92	1.06	2.43	
96	0 87	òòò	85. 0	8.8 E. 6	8.70	8.69	8.69	8.68	8.65	8.60	8.56	8.53	8.42	8.41	8.39	8.39	8.39	8.34	8.31	8.30	8.29	8.25	8.21	8.16	8.15	8.13 8.13	8.08	8.07	8.04	8.03	8.02	8.00	8.00	7.89	7.89	20
,	2.0	9.6	3.32	2.16	39.21	0.80	4.61	9.68	34.28	5.05	0.09	5.11	6.98	11.80	9.51	3.36	0.08	90.0	13.45	10.75	1.09	0.15	1.49	0.16	0.14	21.79	6.11	. 16.15	0.09	0.22	0.08	5.71	8.81	2.29	15.64	
	0.02	0.67	0.38	0.25	4.51	0.09	0.53	1.12	3.96	0.59	0.01	0.60	0.83	1.40	1.13	0.40	0.01	0.01	1.62	1.30	0.13	0.02	0.18	0.05	0.02	2.68	9.76	2.00	0.01	0.03	0.0	0.71	1.10	000	1 98	<u> </u>
:	AA416970	W68220	H41203	H16772	AA629999	AA778646	W72671	AA609471	AA775863	R40373	AA496998	AA682321	N80764	T87515	W47364	AA504265	AA036974	AA132226	N74617	AA488324	R61372	N50806	AA058576	R47893	AA131239	AA447504	AA699427	AA701351	AA460961	AA457576	H94944	AA487674	H51765	A 201173	14/00381	0000
	730036	342640	175533	50227	884511	1048995	345761	743589	878544	28927	897546	462926	300615	115408	324717	825416	484535	566887	296429	842968	37980	280882	380884	153355	503581	784285	433253	435488	796117	838776	230261	841620	103897	100001	020107	770114
	629	099	661	662	663	664	665	999	667	899	699	670	671	672	673	674	675	676	1,1	, K	679	680	8 8	683	683	684	685	686	687	688	689	000	5 6	- 6 6	269	2

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes Upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

7	8	7	7	7	7	7	7	2	7	7	8	8	N	8	74	7	7	8	~	8	8	8	7	7	7	2	8	7	7	8	7	8	8	7		
2.87	3.76	6.01	18.64	0.99	15.20	4.90	9.28	1.00	17.46	9.03	6.84	2.00	3.72	5.66	9.32	15.79	2.32	15.99	88.0	15.83	1.19	9.18	4.60	12.04	9.29	15.98	3.75	8.02	15.57	14.40	10.67	1.23	10.22	12.41		
16.63	1.50	15.21	3.07	4,09	7.43	17.32	12.63	14 11	3.87	11.89	14.72	8.59	16.52	16.02	12.70	5.74	17.38	9 6	12.36	35 Y	18 13	660	15 92	8.97	11.96	4.20	16.45	13.66	4.08		800	48.05	60.0	9000		
4.16	18.23	7.07	1 63	10.1	72.0	70.0	9.5	1.10	7.90	80. 60.	1 41	12.36	2 69	1.23	58.0 88.0	50:5 4 2 7	<u> </u>	3.02	† 46 66 67	0.30	1.23		12.20	1.02	9 6	68.0 68.0		75.0	5 6	S 7.7	6.63	<u> </u>	26.4	1.37	57.	
7 80	9 6	60.7	7.83	97.7	4.74	1.73	7.73	7.70	7.69	7.67	7.07	7.00	6.9	7.04 4.04	, o	10.7	7.60	7.5/	7.54	7.53	7.49	84.7	7.46	7.45	7.44	7.41	7.30	S 1	\$?	7.31 - 25	7.29	7.27	7.27	7.24	7.23	21
c c	0:00 0:00 0:00	7.39	5.20	21.45	1.06	0.14	1.86	2.05	0.08	22.51	0.53	0.25	1.03	23.26	18.06	0.08	5.46	12.06	0.08	7.28	1.18	5.21	0.08	2.44	6.14	0.31	8.77	44.4	9.01	8.19	28.27	15.11	3.80	0.35	2.20	
	- 08	9. 2.	99.0	2.76	0.14	0.02	0.24	0.27	0.01	2.93	0.07	0.03	0.13	9. 40.	2.36	0.01	0.72	1.59	0.01	0.97	0.16	0.70	0.01	0.33	0.83	0.0 Q	1.19	0.60	1.35	1.12	3.88	2.08	0.52	0.05	0.30	
	N62132	AA826324	R50775	W74362	H14949	AA156795	AA453437	H57082	AA463982	AA700739	H10231	H10068	AA780365	AA088886	W72666	T95804	AA291773	R59197	AA702802	R53455	AA633835	AA708441	AA485749	AA702973	R12473	R49644	AA521384	AA011390	N62936	AA459008	T66816	AA706968	R42946	N22766	R43308	
	287598	1420830	38833	345090	48330	502444	788217	204644	810661	435351	46927	46740	1035664	561971	345743	120695	725340	41789	448032	39833	868169	506018	811166	447167	128243	37538	826995	429519	278729	814288	66317	451907	31299	266631	32756	
	694	695	969	697	698	669	700	701	702	703	8	705	706	707	708	709	710	711	712	713	714	715	716	717	718	719	720	721	722	723	724	725	726	727	728	

2 (7	74	7	7	74	8	7	7	8	7	8	7	7	7	64	7	7	8	64	7	8	7	7	8	8	7	7	7	7	7	8	7	7	8		
6.82	6.11	6.43	11.27	15.36	16.05	14.61	10.38	11.80	7.84	7.62	5.32	8.84	0.77	6.81	0.27	4.28	11.87	14.49	3.14	9.27	10.38	15.45	1.16	8.65	12.53	8.19	13.46	9.34	11.61	3.78	5.09	6.48	15.52	3.19	,	
13.06	12.65	13.46	7.61	3.28	3.24	4.57	, 69 60 60 60 60 60 60 60 60 60 60 60 60 60	9 75	5.6 60 60 60	11.28	2.34	11.35	15.66	12.50	17.83	14.80	7.50	5.67	16.93	9.22	8.38	1.38	5.14	9.72	1.10	1.51	2.38	8.48	7.15	2.82	13.88	12.73	. 80 80	16.50	2	
1.81	2.92	4.78	5	7 70 0	2.94	67.7	76.7	0 77.7	7.67	13.51	13.68	1 14	8 8	5 6	2 2	t 80	06. 0	9 6	9 9	0.00	5.5		5.5	25.5	8	10.95	4.74	2 73	, F		5.30	20.1	77.	0 g	8 5	
7.23	7 23	2 6	7.77	02.7	7.19	7.18	7.18	7.17	7.14	7.13	7.12	- ;			11.7	7.05	7.02	7.02	7.00	6.98	6.95	C 70.	0 0 4	ი ი		9 6 8 8 8 8	9.00 98.00 98.00		0. 6. 6. 6.	0. 0 0. 0	6.83	6.83	6.80	6.80	6.79	22
1 17		D	0.31	4.64	11.27	8.56	0.24	15.96	0.11	105.24	14.44	0.62	.51	3.14	95.95	0.26	2.95	2.01	0.28	1.75	0.68	36.26	0.05	0.07	3.78	0.06	2.74	73.80	4.75	0.83	3.35	0.25	2.79	4.70	5.63	
9 46	2 ;	1.24	9.0	0.64	1.57	1.19	0.03	2.23	0.02	14.77	2.03	60.0	0.21	0. 44.	13.50	0.04	0.42	0.29	2	0.25	0.10	5.22	0.01	0.01	0.55	0.01	0.40	3.48	0.69	0.12	0.49	0.0 40	0.41	0.69	0.83	
	H19234	AA135809	AA011182	AA455521	AA431773	AA131909	AA128005	AA504201	AA677655	AA894557	AA455237	AA449118	R96520	AA455087	N66158	AA029722	AA598849	N53940	W72556	AA417288	T68878	AA485453	AA486741	T58146	AA886739	AA669222	AA452278	T63980	AA129135	A1025259	AA935560	AA398406	H56903	R23246	AA455940	
,	50904	565734	359722	809828	782503	504308	501868	825282	897271	1416782	814779	785845	199644	812294	278531	366778	898333	247050	345330	731139	82215	811067	841221	79254	1500162	856585	787938	79761	586650	1641367	1556526	726791	204580	131094	813275	
	729	730	731	732	733	734	735	736	737	738	739	740	741	742	743	744	745	746	747	748	749	750	751	752	753	754	755	756	757	758	759	760	761	762	763	

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell tines appregulated 3 fold or higher in at least 1 of 3 cancer cell lines.

2	7	8	7	7	7	8	7	8	7	7	8	~	~	7	7	7	7	7	7	7	61	~	7	7	7	7	7	7	7	7	7	~	7	7	
1.76	0.98	8.83	3.12	7.33	2.21	13.94	14.72	4.64	4.14	4.12	8.09	1.85	9.54	4.54	4.81	1.28	9.68	6.74	13.83	5.55	6.04 40.04	3.84	2.39	12.67	7.59	8.78	6.08	9.70	1.28	12.22	4.18	8.37	3.99	12.04	
15.10	7.25	1.82	16.41	10.73	14.81	4.25	3.97	14.02	13.65	1.82	0.58	10.05	9.16	12.93	13.90	13.84	7.59	11.33	4.25	11.59	2.65	13.27	3.01	1.84	9.29	9.13	11.14	6.70	12.25	90.9	1.59	7.88	13.18	5.53	
3.45	12.08	9.56	0.64	2.08	3.11	1.87	1.33	1.35	2.21	14.01	11.27	7.97	1.16	2.28	1.02	4.61	2.44	1.47	1.45	2.35	10.79	2.34	14.04	4.90	2.50	1.38	2.05	2.85	5.69	0.90	13.36	2.87	1.95	1.54	
6.77	6.77	6.74	6.73	6.71	6.71	69.9	6.68	6.67	6.67	6.65	6.64	6.62	6.62	6.58	6.58	6.58	6.57	6.51	6.51	6.50	6.49	6.49	6.48	6.47	6.46	6.43	6.42	6.41	6.41	6.39	6.37	6.37	6.37	6.37	23
7.09	1.65	10.08	88.6	7.44	3.87	421.08	3.42	3.96	3.02	17.94	7.75	0.07	63.60	2.64	1.48	1.57	0.12	0.24	5.27	6.55	8.71	10.14	2.76	0.07	0.12	27.63	0.45	6.74	0.21	1.66	9.86	3,76	0.11	4.54	
50.5	0.24	1.50	1.47	-	0.58	82 GR	0.51	0.59	0.45	2.70	1.17	0.0	9.61	0.40	0.22	0.24	0.02	9	0.81	10.	¥.	1.56	0.43	0.0	0.02	4.30	0.07	1.05	0.03	0.26	1.55	0.59	0.02	0.71	
7079VN	AA454562	AA454597	NAZOOS	A A 3 9 8 9 4 9	HORORA	A A B E 7 4 13	PE3448	W37447	44883301	AA165678	AA424790	AA48400	T55728	AA142869	AA029997	H09997	AA039851	T51617	N79745	R43328	AA953357	AA598796	R32406	AA 129677	AA428186	AA626788	AA452542	AA448664	H15910	R89846	N25936	N39611	AA485351	AA405533	
702076	2000a	811582	280154	725677	43750	4475008	147.5045	321900	1461316	593185	768172	781362	73638	504396	470001	46916	375827	72441	300276	32134	1573251	890898	135688	565493	773564	877784	788488	786078	159462	194717	268711	277112	810996	772410	
e e	5 5	76.6	7 2	/0/ 268	90 /	j	i	3 3	1	3 2	7,	7,7	2 12	778	6. 1.	780	781	782	282	78	785	786	787	7 2	280	9 6	3 5	792	793	79.4	705	796	2 6	798	

0 0		7 (7	7	7	7	7	8	7	7	7	7	7	7	8	7	2	2	7	7	7	7	8	7	8	8	7	7	7	8	7	7	7	7		
6.13	C7:11	1.13	11.81	9.86	6.32	8.17	7.26	8.95	7.88	10.08	13.49	5.30	1.69	5.48	90.6	9.66	3.89	6.77	2.74	6.60	7.78	10.29	9.56	9.09	3.85	1.49	7.48	7.07	11.91	5.03	1.00	12.34	12.62	4.79		
12.20	6.38	4.43	4.38	8.03	10.78	8.05	10.13	2.46	8.82	6.21	5.01	0.27	3.53	12.34	8.92	7.60	12.42	0.76	3.73	2.5°	2.03	7.55	2.69	7.76	13.05	12.18	9.73	8.86	3,64	11.73	3.46	4.07	0.62	11.45		
97.0	1.48	13.51	2.86	90	70	5 - C	2.54 4.54	7.40	2 44	2.51	72.0	13.18	13.52	0.87	90.0		 	2.20 44.05	11.03	17.21	1.20	0.40		2 6	5 7	£	- + + + + + + + + + + + + + + + + + + +	3.39	2.75	1.73	13.72	7.5	4 5 4	, ,	<u>3</u>	
6.36	6.36	6.36	6 35	0. c	16.9	6.30	6.29	6.28	6.27	6.27	9.5	0.20 0.20	0.20	0.64	6.23	6.22	6.20	6.20	6.20	6.19	6.16	6.15	6.13	6.13 5.00	6.13	6.12	6.12	- 7	6.11	6.10 00.0	6.03 6.03	9.00	6.03	6.03	6.02	1-7
1.50	6.46	3 38	D	26.80	4.87	51.96	5.87	2.21	1.37	1.58	0.15	2.29	13.69	4.98	4.11	1.36	12.01	3.99	90.0	8.84	0.33	2.57	15.94	12.43	2.89	10.01	0.15	2.40	16.02	207.98	26.54	90.0	0.70	5.33	16.34	
0.24	5	5 5	0.53	4.22	0.77	8.25	0.93	0.35	0.22	0.25	0.02	0.37	2.19	0.80	99.0	0.22	1.94	0.64	0.01	1.43	0.05	0.42	2.60	2.03	0.47	1.63	0.02	0.39	2.62	34.08	4.36	0.01	0.12	0.88	2.71	
89080	000000	AA465223	AA777555	171991	AA459244	AA042812	W85878	N63988	N77828	H54659	R44201	N24869	AA430367	AA877997	AA701978	R38539	H22568	AA630373	W31389	AA779949	N56875	AA865924	AA630104	AA430744	AA609891	AA608752	R26456	R92186	AA430052	AA410517	AA459106	R44617	AA775557	AA165313	AA453441	
70904	120024	814211	448417	85409	814444	486538	416075	289496	302221	203179	34254	267135	769857	1475308	435957	23073	52128	854897	320602	461933	277476	1470195	854701	770992	1032080	950945	133192	195817	781109	753862	814303	33342	378016	594693	788225	
i	799	8	801	802	803	804	805	908	807	808	808	810	811	812	813	814		816	817	. 6	819	820	821	822	823	824	825	826	827	828	829	830	83.1	832	833	

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatio Epithelial Cell Lines appregulated 3 fold or higher in at least 1 of 3 cancer cell lines.

7	7	7	7	7	7	7	7	7	7	7	7	7	7	7	7	7	~	7	7	7	(3	74	7	7	74	7	81	71	74	~	7	~1	2	74	
4.18	5.20	2.68	7.25	7.87	7.79	4.51	11.52	6.52	5.38	60.9	8.76	1.04 4	2.86	4.05	8.22	8.94	4.16	4.04	6.64	2.11	9.69	11.41	5.62	3.29	1.09	12.39	5.26	5,11	4.82	1.55	11.63	8.40	7.45	5.77	
11.73	9.92	10.65	9.75	7.62	8.57	11.00	3.88	9.04	11.46	0:30	7.34	3.44	8.19	11.44	8.58	6.81	12.13	2.57	2.32	12.15	1.00	3.75	9.55	12.93	13.23	4.68	10.89	10.22	11.11	10.45	1.25	7.72	7.30	9.02	
2.14	2.89	4.63	0.97	2.44	1.57	2.42	2.48	2.30	0.97	11.41	1.63	13.24	6.66	2.20	0.84	1.87	1.25	10.90	8.54	3.22	6.78	2.25	2.24	1.15	3.05	0.30	1.21	1.96	1.36	5.27	4.39	1.10	2.46	2.42	
6.02	6.00	5.99	5,99	5.98	5.98	5.98	5.96	5.96	5.93	5.93	5.91	5.91	5.90	5.89	5.88	5.88	5.85	5.84	5.83	5.83	5.82	5.80	5.80	5.79	5.79	5.79	5.79	5.76	5.76	5.76	5.76	5.74	5.74	5.73	52
0.39	7.71	0.54	8.89	8.34	2.99	7.64	0.49	0.55	0.69	1.85	15.84	5.37	0.36	6.55	0.76	2.56	0.30	1.67	4.03	0.41	90.0	7.68	4.83	4.3	4.01	0.14	0.22	0.59	2.01	4.73	0.84	2.62	3.37	6.12	
0.07	1.29	0.09	1.48	1.39	0.50	1 28	80.0	900	0.12	0.31	2.68	0.91	0.0	1.	0.13	0.43	0.05	0.29	0.69	0.07	0.01	1.32	0.83	0.75	69.0	0.03	9,0	0.10	0.35	0.82	0.15	0.46	0.59	1.07	
PKO360	AASAGAS	AA705840	AA465353	A A 504202	N30233	115506 115506	H08210	0.000	AA101173	NG2868	N36008	AA456135	N59866	AA429474	AA778603	N73252	AA425664	AA007529	AA709271	AA044565	H02158	AA708446	AA465090	N69068	AA007516	AA010619	AA609914	AA708605	R46794	N62128	R20639	R13434	R50752	H10788	
38027	813410	1292190	814080	825284	276972	70207	45607	1505	563860	289645	272888	796366	284783	771165	1048949	246549	773367	429376	1343468	488516	150623	506032	814995	287349	429352	430336	1031113	506483	36480	287581	26505	28475	38510	46862	
700	t 200	2 8	3 2		3 8	8 8	3 3	5 6	7 6	3	£ \$	848	748	848	2 0	8 2	85.		853	85.4	, R	8 8	8,47	8,8	859	860	881	882	863	864	865	866	867	868	

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in at least 1 of 3 cancer cell lines.

0.24 0.33 0.03 0.03 1.00 0.51 0.01 0.01 0.05	1.35 1.90 0.15 1.58 2.92 7.88	5.71 5.71 5.68 5.68	2.14 2.14 5.72	12.44	3.07 5.36	0 0
0.33 0.03 0.03 1.00 1.39 0.01 1.19 0.05	1.90 0.15 1.58 2.92 7.88	5.71 5.71 5.68 5.68	1.62 2.14 5.72	9.62	5.36	• •
0.03 0.28 1.00 1.39 0.01 0.01 1.19 0.05	0.15 1.58 5.71 2.92 7.88	5.71 5.68 5.68	2.14 5.72	8.02	>	٧
0.28 1.39 0.51 0.01 0.01 0.14 0.05	1.58 5.71 2.92 7.88 7.38	5.68 5.68	5.72			
1.00 1.39 0.51 0.01 0.01 1.19 0.05	5.71 2.92 7.88 2.35	5.68		9.78	1.55	4 (
0.05 1.39 0.04 1.27 3.28 0.05	2.92 7.88 2.35		1.00	6.51	40.00 40.00	v (
1.39 0.01 0.01 3.28 0.05	7.88	5.68	5.83	9.35	1.87	N (
0.41 0.01 0.14 3.28 1.19	235	5.68	1.17	12.71	6. 10 4. 43	4 6
0.01 0.01 3.28 1.19	2	5.68	1.17	8.12 . 3.0	5 + 1	۰ ،
0.01 3.28 1.19 0.05	7.19	5.67	1.02	08.4	12.25	1 0
0.14 3.28 1.19 0.05	90.0	5.67	3.76	1.00	0.46	. 74
3.28 1.19 0.05	0.81	5.66	3.85	5. c	6.19	7
0.05	18.53	5.65	8.40	1.37	5.00	8
0.05	6.68	5.64	10.55	j.;	, 40 10 10	- 7
	0.25	5.62	0.55	77.7	11.39	8
0.02	0.10	5.62	5.02	; ;	2.59	8
0.01	90.0	5.62		7 70	82.6	8
0.03	0.18	5.61	1.75	94. 4	7.48	. 0
3.17	17.75	5.60	8.06 9.06	07.1	5,99	8
0.01	0.07	5.60	0.80	0.01	5.46	7
0.03	0.17	5.59	1.43	9.00	2.47	8
0.72	4.03	5.59	. 6.89 8.89	4.4	4.42	8
1.97	11.01	5.58	ο <u>ι</u> .:	4.37	4.36	8
0.70	3.89	5.55	11.02	1.27 28.5	29.2	8
99.0	3.65	5.55	0. c	12 B3	0.55	8
0.01	0.05	5.53	5.23 5.00	2	8.52	8
23.98	132.57	5.53	20.1	\$ 2	6.45	. 74
0.07	0.41	5.52	2 7		3.69	8
0.99	5.42	5.50	77.7	5 6	- K	2
12.19	66.94	5.49	7.57))) !	4 6	, 0
2.02	11.11	5.49	0.87	7.74	8. 1	4 (
1.63	8.93	5.48	7.11	7.64	1.70	4 (
0.01	0.05	5.48	3.34	1.00	12.10	N (
88.0	5.38	5.47	0.67	8.14	7.61	81
00.	7 03	5.47	1.85	10.90	3.66	N
0.41	2.26	5.46	4.45	1.50	10.42	0
•		96				

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

7	7	7	7	7	7	7	7	7	7	7	7	7	7	64	8	8	8	7	8	7	7	7	7	7	73	7	7	8	7	7	7	7	7	8	
6.30	7.49	8.44	5.51	8.69	7.71	11.81	3.53	4.35	8.77	4.15	11.07	8.24	4.21	6.03	7.16	7.71	10.03	60.6	5.20	7.18	5.27	6.95	10.80	7.50	7.02	11.04	8.43	6.43	7.68	0.55	1.88	3.74	1.61	6.64	
8.29	6.97	6.79	8.15	5.96	6.69	3.31	10.54	10.79	5.32	10.96	1.00	6.68	10.44	1.01	7.63	7.20	3.73	5.86	9.61	6.76·	8.00	6.68	0.18	2.95	2.97	4.37	5.07	7.32	5.49	10.92	10.57	1.80	10.71	5.99	
1.75	1.86	1.09	2.62	1.61	1.84	1.10	2.12	<u>.</u> 9.	2.06	1.01	4.05	1.13	1.34	8.91	1.15	1.02	2.14	0.94	1.07	1.93	2.59	2.22	4.81	5.35	5.79	0.36	2.21	1.93	2.51	4.20	3.19	10.06	3.27	2.95	
5.45	5.44	5.4	5.43	5.42	5.41	5.40	5.40	5.39	5.38	5,38	5.38	5.35	5.33	5.32	5.31	5.31	5.30	5.30	5.29	5.29	5.29	5.28	5.27	5.26	5.26	5.26	5.24	5.23	5.23	5.22	5.21	5.20	5.20	5.19	27
900	0.05	1.05	0.45	33.51	0.59	001	0.12	2.76	7 13	11 29	0.05	11.61	0.28	0.05	6.28	2.11	14.65	21.66	4.03	0.23	20.12	0.41	4.49	3.68	0.14	0.15	0.58	44.95	15.25	0.08	1.87	0.76	0.46	0.35	
30	3 5	5 6	2 6	8 48		, c	2 6		- C	2 C	2 6	2 47	0.05	0.0	1.48	0.40	2.76	60.4	0.76	9.0	3.81	0.08	0.85	0.70	0.03	0.03	0.11	8.60	2.82	0.02	0.36	0.15	600	0.07	
1	A4426514	190000 00000	A A B B 4 4 7 B	A A 706820	A A 883400	A060184	773034	1,3630	T10030	AA45562/	DA4202	A A 402066	W72310	B73525	D16676	AASORER	AA775865	A A 4 5 5 5 7	N21334	N57858	777840	44628190	VAT2679	14030A	AA879474	NOOSAR	AA620614	AA133166	AA160645	AA404290	AA284280	1872272	00121504	AA446748	
	0/15//	106786	1937.00	464805	4467764	1407343	1300344 9FFB4	0000	96699	004500	34467	744700	245077	156473	129664	808258	878550	813007	265102	247089	108395	1055831	245787	204200	1505735	264646	1048685	490753	504517	758360	324665	2242BG	780884	784126	
	8	2 2 3 3	9 6	3 8		9 6	2 3	5 6	21.6	2 3	2 d	2 6	2 6	, a	5 6	6 C	5 6	, ,	326	924	3,0	920	920	78	078	676	93.5	2 6	33 6	2 6	938	9 6	9 6	88	

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostation Epithelial Cell Line Genes upregulated 3 fold or higher in at least 1 of 3 cancer cell lines.

8	~	7	7	8	7	7	8	8	8	8		۰ ۱	, (4 (v	78 (71	7	8	7	7	7	7	7	8	7	8	8	7	~	8	7	8	0		4	
6.90	5.15	7.33	10.09	7.94	10.01	3.56	8.34	5.28	5.12	7.63	23.0	1 87 Y) i) O.C. 1	7.92	5.79	1.87	3.42	6.61	5.79	76.0	5.65	6.75	6.79	2.27	6.33	6.25	5.46	7.16	10.79	6.59	1.00	5.32	5) (3 8	0.00	
6.91	9.14	1.00	1.15	0.80	3.54	11.06	5.30	8.40	20.6	F 20	0.23	0.02 4	0.12	8.65	5.51	8.28	6.10	10.25	6.33	8.07	5.74	8.14	6.08	6.72	7.27	7.99	6.45	7.62	5.39	3.68	5.47	5.04	8.39	2 2) 60 00 t	5.68	
1.72	1.23	7.12	4.21	6.70	1.82	0.72	168	t		- · c	2.35	6.08 1.08	 40.	0.82	1.63	0.91	7.00	1.25	1.99	0.98	8.10	0.97	1.91	1.20	5.15	0.37	1.97	1.57	2.09	0.13	2.45	8.44		5 1	18.7	2.12	
5.18	5.17	5.15	5,15	5.15	5.12	 	 	- 0	9.10	60.6 1	5.09	5.07	5.05	5.04	5.02	4.99	4.99	4.97	4.97	4.95	4.94	4.92	4.91	4.90	06.4	06.4	9 8 4	88.88	88.88	4.87	4 84	483	9 6	4.00 0	4.80	4.80	78
0.47			22.27	900	800	7 6	0.0	9 (0.43	32.60	1.98	16.55	40.76	0.22	3.45	4.54	8.53	26.53	1.93	6.34	0.05	8.75	0 0 0) ·	 80 0	9 6	0.32	4.49	7 7 7	28 69	10.03	3 2	3 9	13.99	0.05	92.90	
4	3 5	5 5	5 6	4.32	5 6	4 5	P.0.0	0.0 4	0.08	6.40	0.39	3.26	8.07	0.04	0.69	0.91	1.7	5.33	0.39	28	5	1 78) e	3 6	? 6	7 0.0	2 7	r 6		- 6	0.90	2 7	0.0	2.91	0.01	19.35	
	AA644334	AA426361	AA398218	AA129931	W84774	H99997	R38169	AA683338	AA777779	AA148505	R71689	AA962541	AA421230	AA490605	N49899	H61552	AA149637	AA504631	N20480	D44346	N44340	138200 4 4 4 4 5 6 4 3	AA4/0342	K82041	78698N	AA114106	H16743	AA804808	R33/30	AA035384	AAU43228	NO23/2	AA454554	AA064627	R43271	AA456931	
6	845002	708140	720038	501934	415806	263342	23658	1293016	448920	491524	155575	1602209	739247	824109	243653	206052	504279	825603	264162	34704	406432	201001	711401	14/925	277545	563201	50114	1504152	12/04	4/1098	485/8/	000087	803208	382451	32517	838568	
,	939	940	<u>4</u>	942	943	8	945	946	947	948	949	950	951	952	953	95	95	9 4	2 2	200	0 0	n (98	961	962	863	964	965	998	296	896	9	970	971	972	973	

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line. Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

7	7	7	7	7	7	7	7	7	7	7	8	7	7	7	7	7	7	7	7	7	7	7	7	8	7	7	7	7	7	7	7	7	8	74	
7.20	5.62	0.65	5.93	6.25	6.47	6.37	8.48	6.21	6.12	5.80	1.44	5.38	6:39	5.88	6.25	5.98	5.05	7.00	2.28	5.85	5.23	5.23	7.00	5.54	5.12	5.56	6.95	5.02	5.57	6.78	1.71	5.93	5.98	5.87	
5.26	6.24	10.08	6.79	6.82	6.47	5.38	0.08	5.74	5.51	1.33	5.90	2.63	5.33	5.63	5.20	5.24	5.72	0.75	5.06	6.62	7.47	7.39	5.53	5.84	6.30	2.55	5.28	6.17	6.98	5.09	5.25	6.24	5.02	5.71	
1.94	2.51	3.62	1.60	1.18	1.30	2.49	5.66	2.27	2.52	6.88	6.62	5.94	2.19	2.36	2.32	2.55	2.99	5.99	6.34	1.15	0.93	0.99	1.08	2.21	2.12	5.38	1.27	2.26	0.90	1.56	6.45	1.21	2.31	1.72	-
4.80	4.79	4.78	4.77	4.75	4.75	4.75	4.74	4.74	4.72	4.67	4.65	4.65	4.63	4.62	4.59	4.59	4.59	4.58	4.56	4.54	4.54	4.54	4.54	4.53	4.52	4.50	4.50	4.49	4.48	4.48	4.47	4.46	4.44	4.44	59
7.55	24.67	0.10	2 6	. 4	090	4.36	16.44	0.34	4 54	0.12	0.27	1.82	4.18	5.17	15.49	4,45	4.50	90.0	12.03	13.59	5.05	2.67	0.38	2.96	40.40	2.71	6.39	4.53	1,60	0.32	1.09	0.53	0.06	0.49	
1 57	. Y		2, 6	, c	5 6	2 6	2 47	; e	3 6	8 6	800	0.39	0.90	1.12	3.37	0.97	0.98	0.0	2.64	2.99	1.1	0.59	0.08	0.65	8.95	0.60	1.42	1.01	0.36	0.07	0.24	0.12	0.01	0.11	
1134614	101041	AX131320	45477	704640	1000010 444444	AA321411	K002/3	AA//9103	74/00413	00/00	AA676590	H08796	AA443094	AA626847	AA598797	AA460521	AA284291	AA629251	AA676296	H98655	AA488979	N51367	AA704858	R24451	AA421270	AA885096	H58542	T58434	N29624	AA778562	H22854	AA883660	H20743	AA904776	
90090	008007	2027.13	20040	194472	671407	800070	667671	455000	40000	43020	432732 882488	46097	809466	745192	898083	796606	327245	744044	882355	261567	824728	283208	453229	132954	731031	1468466	206272	75650	257312	1048899	51700	1466599	51328	1504005	,
į	d 1	6/6	9/6	977	9/6	D (0	9	- SG	706	66.0	4 W	0) k	888	8 0	8 8	8 6	. 6	* 60 0	766	, c	966	266	855	666	1000	1001	1002	1003	1001	1005	100	1007	1008	

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line Centes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

7	7	7	8	7	73	74	8	8	8	- 6	. 7	7	7	8	8	۰ ۱	4 (ν (N (N (N	7	7	7	7	7	7	7	7	7	7	8	2	7		•	
6.76	5.17	5.29	5.41	6.30	5.23	5.54	5.26	5.86	470	6.47	58.84 44.	5.62	5 79	6.07	36	, i	0 10	5.87	5.58	5.36	5.41	5.49	5.65	6.01	5.93	5.55	1.12	5.01	0.56	5.56	5.46	0.55	5.01	563	2.63	} •	
5.63	2.84	0.22	5.60	0.31	5.46	5.14	: Z	, r	90 90		. G.	5.00 7.00	5,5		. 4	\$ i	5.55	1.07	5.50	5.73	1.63	6.05	5.14	5.73	5.12	5.71	5.23	1.78	5.82	5.04	5.12	5.69	5.60) u	9.00	3000.91	
0.87	5.24	7.74	2.19	E 57	2.44	9 6	2.30	4 · · ·	78.1	5.57	9 0	1.30		97.1	90:-	1.91	1.25	5.58	1.40	1.38	5.38	0.84	1.58	0.57	1.23	0.97	5.88	5.25	5.37	1.13	1.12	5.46	2 7	5. 6	5 to 1	0,55	
4.42	4.42	4.42	7.40		. t.	8.4 8 r	6.35	4.35	4. 4.	4.33	4.32	4.32	4.28	4.28	4.25	4.24	4.19	4.17	4.16	4.16	4.14	4.13	4.12	4.10	4.09	4 08	4 08 80	4.01	3 92	3.91	3.90	00 %	9 6	3.87	3.87	1221.36	30
1.88	800	1 0	7:7	57.53	41.0	0.84	0.73	0.40	0.42	0.10	1.07	22.74	0.39	0.27	6.07	11.37	8.82	5.73	11.43	0.36	1.69	9.76	0.57	0.78	0 :: C		 	C2.4	20.0	5.5	5.6	6.0	5 ·	1.56	7.56	12.21	
0.42		0.02	1.71	13.07	0.03	0.19	0.17	60.0	0.10	0.02	0.25	5.27	0.09	90:0	1.43	2.68	2.10	1.37	2.75	60.0	0.41	- 5.6	2 7	† °	9 6	0.50	7. 7	<u>.</u>	20:- 0	0.02	77.7	0 (6	0.02	0.40	1.95	0.01	
301.000	R09/00	R06372	AA398355	T90621	AA404338	AA112979	AA884382	AA626247	AA625791	AA443300	AA205320	T70122	AA437245	H08227	N50828	AA465692	N35579	R60949	NEOGRA	A A B 8 3 7 5 5	A A 708 424	AA 187206	AA407.200	K0/012	AA633993	AA862814	154527	162552	N/8230	H79566	AA676598	H84915	N58136	H47089	AA045340	N32071	
770707	1040	126239	726858	110772	758302	530545	1461715	745523	744911	784589	647816	80946	757463	45514	280909	814906	272238	42400	284103	4466824	130001	200987	841287	126695	858292	146921	73659	79726	295787	239661	882483	249517	247587	193420	487141	260168	
•	1009	010	101	1012	1013	1014	1015	1016	1017	1018	1019	1020	1021	1022	1023	1024	1025	400	020	707	070	9201	1030	133	1032	1033	1034 4	1035	1036	1037	1038	1039	5 5 6	1941	1042	1043	

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostate Epithelial Cell Line: Defice upregulated 3 fold or higher in at least 1 of 3 cancer cell lines.

-	~	-	-	-	-		-	y-	-	-	-	-	-	-	~	-	•	-	*	-	-	-	-	•	₩-	-	₩-	-	-		-	~~	-	-	
1.00	0.59	1.33	1.09	1.00	2.22	2.96	2.07	0.74	2.83	1.09	 2	0.74	0.30	2.47	1.77	0.29	97.0	0.38	. 2	1.00	0.97	1.10	0.36	2.35	1.26	2.37	0.68	69.0	0.07	0.48	1.06	1.96	1.21	0.19	
1675.21	1262.88	1231.52	761.20	730.07	708.49	652.56	586.88	515.39	458.06	446.45	438.73	264.42	241.37	221.03	220.07	218.92	218.05	214.82	212.83	191.25	176.58	175.32	173.57	168.03	162.70	161.00	153.57	149.67	151.46	148.64	143.96	135.65	135.09	137.23	
1.63	1.71	0.98	1.49	2.76	2.72	2.43	2.91	0.55	1.62	0.82	1.13	1.24	0.85	1.50	1.79	0.77	0.60	2.48	2.70	1.54	1.07	1.25	1.85	2.01	1.13	0.86	0.97	2.74	0.39	1.48	0.82	2.69	3.00	0.95	
559.28	421.73	411.27	254.59	244.61	237.81	219.32	197.29	172.23	154.17	149.45	146.97	88.80	80.84	75.00	74.54	73.32	73.13	72.56	72.19	64.60	59.54	59.22	58.59	57.46	55.03	54.74	51.74	51.03	50.64	50.20	48.61	46.77	46.43	46.12	31
5.59	7.11	158.35	2.55	2.45	2.38	2.19	1.97	6.87	6.56	73.56	49.37	69.16	7.42	3.61	8.93	16.59	23.96	1.43	2.83	0.65	3.72	20.73	18.15	1.84	14.22	3.65	4.65	0.94	7.71	1.05	9.24	32.83	1.25	12.55	
000	0.02	0.39	0.01	0.01	0.01	0.01	0.0	9.0	9.0	0.49	0.34	0.78	0.09	0.05	0.12	0.23	0.33	0.02	9.0	0.01	90.0	0.35	0.31	0.03	0.26	0.07	0.09	0.02	0.15	0.02	0.19	0.70	0.03	0.27	
NA7064	N93470	N48913	N30131	R01937	W20486	AA476502	AA486538	AA875888	AA432058	AA456130	AA406070	AA401342	AA461320	AA426212	H11642	AA598983	N33323	AA399245	H05961	R61395	AA133554	AA002166	AA171784	H16537	AA432278	AA459400	AA481806	AA001219	AA621324	H47076	AA013353	AA398384	AA443140	AA456951	
280204	307328	279613	268115	124567	327495	785337	840937	1325605	784116	796350	743027	742564	796334	769542	47963	897722	270603	726439	44007	37986	586845	427693	594683	49131	782353	810959	838877	362278	1048794	193087	360355	726934	796723	838600	
770	1045	1046	1047	1048	1049	1050	1051	1052	1053	1054	1055	1056	1057	1058	1059	1060	1061	1062	1063	1064	1065	1066	1067	1068	1069	1070	1071	1072	1073	1074	1075	1076	1077	1078	

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least for 3 bancer cell lines.

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0.57	2.63	0.54	3 3 3		1.42	0.44	2.13	1.07	2.03	0.87	0.13	1.83	9.84	0.39	0.18	108.31	55	. t	- c	2.00	1.78	2.6	95.10	2.37	0.86	1.00	0.39	1.00	0.25	2.69	0.76	1.56	0.82	2.73	34 5	2	<u>8</u>	
130.86	128.29	120 35	129.55	122.8/	1.33	122.05	120.12	1.22	117.01	117.08	500	24 50	413.87	- + 	10.30	5	1.00	106.62	106.86	104.16	99.56	98.13	1.00	90.43	92.44	90.70	91.71	84.23	82.35	79.62	78.91	79 11	0 × 0 ×	1 000	(0.73	1.76	76.96	
1.23	2 6	8 · ·	1.18	0.47	122.01	0.72	0.41	11816	5	3 4	1.18	718.3/	0 4	CL.1	0.83) (0.73	0.68	0.12	1.55	1.05	1.40	1.00	1.86	1 16	2.74		55.	2 · ·	; c		- 2:7	2 7	1.24	1.91	1.43	1.74	
44.22	77.55	43.90	43.69	41.89	41.59	41.07	40.88	50.04	40.15	40.03	39.71	39.51	39.20	38.61	37.26	36.82	36.69	36.32	36.21	35.93	34.13	33.51	32.37	31.55	2. 2. 2. 2. 2. 2. 2. 2. 2. 2. 2. 2. 2. 2	31.40	04.10	31.06	28.99	16.82	28.14	27.29	27.26	27.13	27.12	26.76	26.59	32
ļ	0.77	1.29	21.09	0.63	4.73	42.43	C#:7#	4. 4.	15.59	1.72	23.11	14.33	10.92	30.27	82.43	2.07	13.35	16.34	1.94	10.68	13.40	75.0	33	20.0	0.32	37.35	0.31	1.21	5.31	<u>.</u> 26.	0.63	0.36	25.14	5.54	1.91	12.79	2.37	
	0.05	0.03	0.48	0.02		- 6	50.F	0.10	0.39	9.0	0.58	0.36	0.28	0.78	2.21	90.0	0.36	0.45	0.05	030	95.0	6 6	5.0	0.01	0.01	1.19 6	0.01	9.0	0.18	90.0	0.02	0.01	0.92	0.20	0.07	0.48	60'0) }
	W70264	N34482	AA25743	A A 4 9 7 5 0 5	000000	H08850	W86779	W42996	N64840	N27641	N26083	H23524	W88720	AA180453	AA481788	N34943	AA677643	AA457138	D10140	20000	N23003	N40040	N52158	H57136	AA449300	N48271	AA159497	AA128407	AA704816	W43028	AA121387	AA872602	N66454	R37289	AA433910	A A R 73604	AA447764	Www. 1. 2.
	344834	271141	272200	2010	010070	46055	416808	323251	284701	255659	268979	51799	417760	609436	838853	276905	897262	810459	428484	104071	294343	9/18/2	284408	204686	785866	282161	592403	526567	452848	329059	564231	1323328	285581	27152	773678	132348	043664	0.000
	1079	080	200	9	1082	1083	108 4	1085	1086	1087	1088	1089	0601	100	1092	1093	7004	# NOC +	2000	9601	1097	1098	1099	1100	1101	1102	1103	<u>+</u>	1105	1106	1107	1108	1109	1110			2111	1113

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Lines (Genes upregulated 3 fold or higher in at least 7 of 3 cancer cell lines.

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2.63	1.55	2.80	1,17	0.80	168	2.46) } }	2.00	2.30	1.42	1.00	2.23	2.18	0.64	1.17	1.17	2.77	 	· ·	//·o	0.10	0.87	0.97	2.57	1.38	1.40	0.63	0.68	1.70	59.09	0.10		60 60	0.04	2.16	0.46	0.59	
75.29	73,65	73.28	74.73	74.38	7. CF	##:7/	70.67	71.10	71.45	71.41	70.65	71.20	68.08	69.82	68.34	67.64	64.67	70.40	65.48	66.20	65.41	64.81	64.73	63.34	64.28	62.07	63.23	62.26	59 72	3000	6.06 6.06	90.04	58.64	59.31	56.55	57.49	57.17	
1.00	2.60	131	17:1	n (0.29	1.28	1.92	1.38	0.41	1.19	2.32	0.29	2.54	101	4 52		07.1	1.35	1.39	0.83	1.64 46.1	1.16	0.97	0.70	984	. 55) T	0 H) () ()	[7]	1.29	1.10	1.33	0.81	0.96	1.44	1.24	
26.31	25 93	25.52	0/:07	25.55	25.16	25.13	25.02	24.82	24.72	24.67	24.66	24.57	74.27	2 82	70.07	23.00	23.36	22.90	22.78	22.60	22.38	22.28	22 23	2 2	22.27	24.60	51.69	21.68	21.17	20.87	20.81	20.63	20.35	20.32	19.89	19.79	19.67	33
25 73	27.72	55.45	0.26	3.89	2.31	9.24	4.63	5.49	59.39	16.06	0.25	200		10.63	37.42	10.35	12.38	27.72	6.49	9.38	2.19	. Y	4 100	0.71	18.0	24.0 D	4. 05	12.75	29.99	10.68	0.18	1.66	3.00	5.45	5.04	12.57	11.90	
8	20.1	1.29	0.01	0.15	0.09	0.37	0.18	0.22	2.40) u	3 3	5 6		44.0	1.57	0.44	0.53	1.21	0.28	0.42	5 6	2.0	5 (),'O	0.0 4	0.38	0.19	0.59	1,42	0.51	0.01	0.08	0.15	0.27	0.25	19 0	080	3
	AA776041	N74889	R40057	AA418028	AA425214	AA417280	H15050	040462	701400	000/80	1/6/90	4//40N	AA432096	AA884321	R09504	AA011570	AA022668	AA426030	AA417211	A A 400264	A4480204	K92512	AA406220	N66346	W45403	N36389	W74696	AA470073	H89563	N62079	AA136049	R49436	R19478	R59304	AA885141	NIDORK4	1429001	N9 187
	970901	299603	27544	767739	773293	731115	18363	0000	20200	108107	285682	284569	784154	1466893	127711	429660	364454	757235	731154	902000	08/670	1964/5	742890	285364	328745	273024	344747	730377	253534	289865	502593	38244	33045	37901	1466551	76086	20000	40004
	1114	1115	1116	1117	1118	110		2 3	וצוו	1122	1123	1124	1125	1126	1127	1128	1129	1130	7	2 3	1132	1133	134	1135	1136	1137	1138	1139	1140	1141	1142	1143	1144	1145	2 4	5 1	/#[[1148

Table 2-10; Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Lines Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

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1.00	 64 90	5.20	0.76	0.38	0.41	40.	141	23	1.83	0.50	0.53	1.45	20.	2.29	90	3 6	2 6	n (1.59	2.06	1.55	0.77	0.74	1.16	0.53	1.30	2.52	44.37	2.06	1.23	1 07	. a	0.30	6/.0	0.97	
55.61	53.78	1.00	55.52	7. 2.	£3 43	25.15	01.70	51.00	- 7.7	19:00	50.80	0.87	48 89	16.93	40.90 40.90	49.51	06.1	0.05	46.45	45.77	46.79	47.42	46.41	46.01	46.29	44.76	42.94	00	42 61	5 5	2007	42.03	04.40	43.90	42.90	
0.1	1.56	1.10	2.52	0.98	0.30	F. 0.	1.27	0.55	50.03 10.03	1.35	2.13	 	49.43	0.80	88.	0.36	45.76	48.82	1.51	1.72	0.92	1.01	1.38	1 23	7				20.1	D (0.10 0.10	2.79	1.07	1.08	1.48	
19.20	19.11	19.00	18.85	18.68	18.29	18.05	18.02	17.94	17.66	17.52	17.44	17.28	17.25	17.14	17.05	16.83	16.65	16.52	16.52	16.51	16.42	16.40		, c	0.0	15.07	15.67	10.08	15.56	15.49	15.37	15.30	15.28	15.26	15.12	34
0.19	8.40	0.19	0.19	12.10	0.48	14.96	2.32	17.51	7.11	6.75	1.37	3.05	8.13	1.84	6.13	1.60	5.82	3.57	6.03	5.21	3.70	, ;; ;; ;;	77.1.	85.4 8	14.63	1.63	4.23	5.43	0.16	2.75	8.40	7.61	3.82	8.56	12.92	
0.01	0.44	0.01	0.01	0.65	0.03	0.83	0.13	0.98	0.40	0.39	0.08	0.18	0.47	0.11	0.36	0.10	0.35	0.22	0.37	2		3, 6	9.68	0.31	0.91	0.10	0.27	0.35	0.01	0.18	0.55	0.50	0.25	0.56	0.85	; ;
AA065042	AA167120	W74725	AA421256	T80832	R43189	AA884397	AA725397	H99075	AA634379	AA620607	AA188710	AA625641	AA418000	AA706935	R60731	R43017	AA884926	NSORRO	A A 412403	74416403	BOYLON.	H986/6	AA620301	R68409	N40202	AA190825	R51506	AA495938	N75473	AI005321	N66348	W44338	AA845168	AA421170	AA083514	
525478	603209	344618	731016	109275	32551	1461725	1342650	262060	743828	1048694	626186	745330	767449	451871	42225	31869	1468260	284003	434433	774107	2018/2	261989	1030808	137853	276438	627401	38677	768445	299162	1626197	285370	323599	1412504	731001	£404£7	2
1149	1150	1151	1152	1153	1154	1155	1156	1157	1158	1159	1160	1161	1162	1163	1164	1165	4 5	3 5	7911	2911	1169	170	1171	1172	1173	1174	1175	1176	1177	1178	1179	1180	7 7 7	2 ?	701	3

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in at least 1 of 3 cancer cell lines.

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0.94	1.13	1.29	0.68	1.92	0.42	0.95	1.16	0.08	0.67	38.45	1.48	0.99	1.53	1.31	36.52	1.27	0.85	0.39	2.87	0.71	0.25	1.42	0.57	0.57	0.79	0.65	2.46	1.29	0.72	0.53	0.11	1.60	0.65	1.51	
43.54	42.32	41.82	42.02	40.40	41.08	41.04	40.96	41.37	41.22	1.37	39.26	0.11	39.18	39.02	2.34	39.01	39.43	39.92	36.40	38.96	39.39	38.18	38.68	38.10	38.03	38.31	35.14	37.10	38.34	37.93	38.93	35.65	36.87	36.05	
0.85	1.04	1.17	1.26	1.33	2.10	1.51	0.98	1.45	0.76	2.13	1.19	40.60	0.91	1.22	2.64	1.10	0.91	0.69	1.51	0.89	0.86	0.89	0.86	1.42	1.22	0.95	2.26	1.42	0.73	1.19	0.42	1.76	1.47	1.33	
15.11	14.83	14.76	14.65	14.55	14.54	14.50	14.36	14.30	14.22	13.99	13.98	13.90	13.87	13.85	13.84	13.79	13.73	13.67	13.59	13.52	13.50	13.50	13.37	13.37	13.35	13.30	13.29	13.27	13.26	13.22	13.15	13.01	12.99	12.96	32
5.05	7.34	4.03	8.55	7.71	0.34	21.55	0.78	1.87	8.37	0.14	2.05	1.31	8.10	10.61	0.14	9.64	14.87	2.31	14.75	8.43	2.77	7.69	12.65	3.60	12.03	13.46	0.26	6.68	7.81	4.53	1.19	2.89	14.04	7.09	
0.33	0.50	0.27	0.58	0.53	0.02	1,49	0.05	0.13	0.59	0.0	0.15	0.09	0.58	0.77	0.01	0.70	1.08	0.17	1.09	0.62	0.21	0.57	0.95	0.27	0.90	1.0	0.02	0.50	0.59	0.34	0.09	0.22	1.08	0.55	
AA844124	AA412435	B41782	N48302	N47954	AA911971	T64323	AA211483	AA703660	AA46882	R36969	R61189	H08397	AA463483	N66724	AA621218	W45330	W60894	R61556	AA464541	H99722	AA083207	AA181723	AA417031	R45114	N38992	AA469939	AA158211	AA417017	N51290	AA460967	AA143467	N92310	AA400273	AA609887	
1388373	731444	30986	279616	280270	1474149	80242		450375										38015	810511	262968	549035	612782	730888	34901	243410	730398	592778	730827	283045	796130	591814	308031	742666	1032072	
787	184	2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2	1 2 2		2 5	1180	19	1192	1193	192	14.05	198	1197	1198	1199	1200	1201	1202	1203	1204	1205	1206	1207	1208	1209	1210	121	1212	1213	1214	1215	1216	1217	1218	

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1.62	0.0	0.65	1.30	1,58	1.39	1.00	1.26	2.43		0.76	62.0	35.74	466	 28.0	1.50	1.46	1.28	290		34.33	67:-	0:00	0.80	1.06	1.61	1.68	0.95	1.66	0.40	0.27	1.51	0.90	1.13	0.32	1.42		
36.14	36.21	36.50	1.48	35.01	35.31	35.57	35.21	4.00	03:50	2.55	2) '	8 5	24.01 20.02	33.50	3. C.	32.02	33.72	54.50	2.56	33.75	33.81	32.40	32.62	32.51	31.51	0.28	1.22	32.53	33.21	30.58	31.68	29.73	0.83	30.72		
0.97	1.34	1.20	35.38	1.20	2	5 5	9 9	8 I	1.77	6.5	35.07	z. 2	0.66	1.20	0.77	04.6	2.07	- -	0.97	9.	0.87	0.82	1.91	1.22	0.78	1.68	33.19	31.50	7 30	04: 64:	6 6 7	5 6	0.50	\$ 2	0,20 1,35	<u> </u>	
12.91	12.84	12.78	42.72	12.72	12.60	12.30	12.52	12.49	12.49	12.47	12.44	12.37	12.35	12.29	12.20	12.16	12.10	12.04	11.98	11.96	11.96	11.73	11.70	11.63	11.63	11 62	11.47	11.46	P 6	11.50	11.35	11.26	11.20	11.23	11.23	11.16	36
8.15	99.9	13 65	8.5	02.4	8.45	5.81	0.13	2.68	6.06	8.34	3.96	6.39	0.19	2.21	10.12	5.89	21.58	7.77	7.93	0.12	9.57	10.22	15.85	080	336) c	6.0	2.01	73.54	3.38	2.03	0.41	7.40	0.68	4.08	5.33	
0.63	0.62	0.92	7.0	0.33	0.67	0.46	0.0	0.21	0.49	0.67	0.32	0.52	0.02	0.18	0.83	0.48	1.78	0.65	99.0	0.0	0.80	0.87	. t	9 6	8 6	0.23	0.30	0.23	6.42	0.30	0.18	9. 20.	99.0	90.0	0.36	0.48	
W84815	Cloton	W94591	T64192	AA677165	AA418988	N75394	AA053239	R56642	N66203	AA884317	H62162	AA431750	N74623	AA088701	W92278	AA029428	AA135222	AA400281	A A 6.09138	D17096	000117	A440306	AA 100333	N62700	H55955	AA449745	W74471	AA862465	AA700604	T91244	AA411682	N21233	AA448672	H92234	AA702640	AA469922	
445820	070014	358046	80186	454317	768111	299024	510136	41300	278629	1466883	208413	782279	296448	511233	359040	366795	588159	742671	1031375	420862	143004	180510	623084	788284	203488	785788	346615	1456160	433350	112559	753368	264449	786073	221828	384015	730344	
9	1219	1220	1221	1222	1223	1224	1225	1226	700	1228	2 000	1230	1234	4232	1233	1234	1235	1236		/53/	1238	1239	1240	1241	1242	1243	1244	1245	1246	1247	1248	1249	1250	1251	1252	1253	:

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in at least 1 of 3 cancer cell lines.

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1.05	0.99	0.52	0.76	0.63	0.36	0.67	2.58	0.98	0.84	1.80	1.25	1.56	<u>.</u> ¥	29.91	60.0	0.97	0.24	1.26	0.56	2.05	1.60	1.20	1.00	1.07	0.45	1.21	0.82	1.26	1.36	0.90	1.89	0.88	2.33	0.90	
31.28	30.94	31.46	31.48	31.25	31.52	31.48	29.45	1.01	30.86	29.68	29.36	29.74	29.70	90.0	31.04	29.17	30.56	29.08	29.40	28.01	28.15	28.46	28.58	28.32	29.62	28.06	28.36	27.58	26.99	27.67	26.24	28.25	26.16	27.33	
1.1	1.32	1.26	0.91	1.23	1.13	0.83	0.80	30.68	0.97	0.81	1.66	0.92	1.06	2.06	0.73	1.65	0.75	1.01	1.26	1.13	1.20	1.12	1.20	1.33	0.52	1.31	1.38	1.20	1.65	1.33	1.63	0.49	1.03	1.30	
11.15	11.09	11.08	11.05	11.04	11.00	10.99	10.95	10.89	10.89	10.76	10.75	10.74	10.70	10.68	10.62	10.60	10.52	10.45	10.41	10.40	10.32	10.26	10.26	10.24	10.20	10.20	10.19	10.01	10.00	9.97	9.92	9.87	9.84	9.84	37
9.68	0.79	11.98	7.28	3.51	1.56	3.26	2.62	15.93	9.08	5.00	1.34	5.45	2.56	2.63	1.13	4.27	2.53	4.74	6.94	18.36	3.36	4.50	0.10	3.44	1.53	4.33	14.09	5.32	3.47	6.03	7.21	4.57	6.60	4.44	
0.87	0.07	1.08	0.66	0.32	0.14	0.30	0.24	1.46	0.83	0.46	0.12	0.51	0.24	0.25	0.11	0.40	0.24	0.45	0.67	1.71	0.33	0.44	0.01	9.34	0.15	0.42	1.38	0.53	0.35	0.60	0.73	0.46	0.67	0.45	
Nacror	AA191495	AA188999	A 4 40 60 40	N33550	B40244	H17038	R42763	T68445	N59278	H99460	R41389	N47214	AA463463	H06525	R51236	AA401482	AA219315	R45402	AA236561	AA446658	AA412512	AA885140	T50397	AA460289	AA426038	AA453466	N51107	AA609415	AA598943	AA024459	H06380	H96738	W70242	AA459652	
258063	627292	626462	82361B	270274	28774	50582	31478	83358	289725	262313	29251	280413	811785	44387	38517	742616	629968	35308	687579	784177	730294	1466549	72663	795723	773423	795170	282000	743519	898038	364950	44303	251685	344010	795540	
1361	125.	1256	1257	10.50	4250	1280	1261	1262	1263	1264	1265	1266	1267	1268	1269	1270	1271	1272	1273	1274	1275	1276	1277	1278	1279	1280	1281	1282	1283	1284	1285	1286	1287	1288	

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

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0.95		5	0.73	1.08	26.83	1.31	0.24	1.05	0.43	000	103	23.15	0.93	0.74	2.61	2.67	0.12	25.05	2 88		99.0) -	- c	3 6	; ; ; ;	1 86	1.78	1.78	• •		9 6	6 · · ·	0.0	2.72	1.26	
27.36	27.43	26.98	27.37	27.09	0.40	26.66	79.76	26.48	26.95	20.33	24.40 26.65	2 84	2.50	26.29	23.30	22.00	22.07	77:17	8.5	24.01	25.00	67.67	25.60	25.56	24.09	24.82	2.38	2.30	53.39	24.00	24.09	24.14	25.33	23.64	24.08	
1.16	1.15	1.25	1.14	1.02	1 83	5		4 6	1.51	1.25	2.24	0.87	2.30	74.07		2.22	1.48	0.68	1.90	0.96	1.03	1.55	1.00	1.38	1.09	1.18 1.18	70,1	22.82	1.61	4. 4.	1.57	0.96	0.81	0.35	1.26	
9.82	9.80	9.76	9.75	9 7 3	0 0 0	n (0)	0.6	9.65	9.61	9.56	9.55	9.52	9.51	9.40	9.40	9.38	9.3 4	9.34	9.32	9.28	9.28	9.26	9.22	9.20	9.14	9.07	00.6	8.99	8.99	8.97	8.96	8.96	8.93	8.91	8.87	38
12.27	5.41	5.49	7.41	96 7	0.1	1.71	1.53	1.56	6.78	4.97	9.52	4.77	33.20	3.37	6.72	0.72	1.69	0.94	60:0	7.66	1.40	0.19	0.09	3.06	2.38	3.63	3.37	19.69	0.91	6.33	0.48	3.83	8.93	0.87	0.07	
1.25	0.55	556	9 0	0 0	0.50	0.18	0.16	0.16	0.71	0.52	1.00	0.50	3.49	0.36	0.71	0.08	0.18	0.10	0.0	0.82	0.15	0.05	0.0	0.33	0.26	0.40	0.37	2.19	0.10	0.71	0.05	0.43	1.00	0.10	5	
197599	74770409	VV 9498	AA8/3009	AA406213	N52876	H11730	H52446	H05626	H89331	AA432141	AA199717	R61871	AA115076	AA459941	N70654	H23555	R39901	H05741	771042	AA127069	AA705858	W69649	T49325	W74633	AA774645	N35603	R41357	AA426113	AA485429	N27272	AA490058	W70074	44437107	AA46873	A 5 0 2 5 4 4 4	2000
121633	247000	030/46	1325/51	742874	283619	47793	201931	43422	253314	781506	647397	43072	491565	796398	299347	51906	26844	43815	84141	502664	1292207	343871	67397	346642	399319	272288	29424	757265	811050	257608	839903	343977	757352	F07677	30202	200
980	807	1290	1291	1292	1293	1294	1295	1296	1297	1298	1299	1300	1301	1302	1303	1304	1305	1306	1307	200	1309	1310	131	1212	4 6 6	1314	1315	1316	1317	1318	1319	12.00	3 5	1351	1322	1323

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line Cent. upregulated 3 fold or higher in at least 1 of 3 cancer cell lines.

	_	-		-	-	-	-		-		-	-	₩-	-		-	-	-	-	-	-	-	-	-	-	Ψ-		-	-	-	-	•	- -	-	
1.47	2.29	1.83	0.13	25.13	2.36	0.95	24.14	2.05	0.67	1.55	2.22	24.73	2.79	1.38	23.39	0.94	0.70	2.24	2.88	0.62	0.97	0.65	20.37	0.47	0.94	0.50	2.96	1.19	1.22	0.80	1.71	0.62	1.74	21.36	
23.71	21.71	2.00	25.76	0.02	21.36	24.92	0.53	23.05	24.06	23.01	21.50	0.28	20.01	22.84	1.42	22.84	23.77	22.41	20.90	23.97	23.00	23.12	2.77	24.31	23.22	24.02	19.19	22.37	22.28	23.03	21.45	1.81	21.62	0.15	
1.28	2.26	22.34	0.22	5 6	233	0 13	1.27	0.83	08.0	00	1.79	0.49	2.69	121	0.57	1.56	9 6 8 6	0.0	2. c		5 4	3. +	3 8	40.0	0.78	35.0	2.68	1 18	4.	080	1.45	22.47	1 13	2.95	
8.82	8 75	9 6.5	9.70	0.70	0.09	Ø.08	0.0	0.00	5 6	, 0, 0 4, 0, 1	0.32	0 0) () ()	0.00	0.40 0.40	0.40 0.40		4. 0 4 0	8.43	8.41	8.40	8.39	8.38	8.38 8.38	8. v	8.5 9.5	67.0 92.0	0.50	9.24	8.21	8.21	8.20	8.20	6.10 0.10	39
75.0	6.5	2.31	10.67	11.41	2.00	0.78	0.78	30.73	3.60	3.84 4	2.73	0.89	0.30	37.45	1.91	8.81	3.53	3.73	20.33	2.92	17.03	2.03	1,94	18.69	3.93	1.42	2.74	10.19	2.38	4.33	8.62	1.81	2.30	3.42	10.88
č	9. 9.	0.26	1.22	1.31	0.58	0.09	60.0	3.55	0.42	0.45	0.32	0.11	9. 2.	4.41	0.23	. 4	0.42	0. 44.	2.41	0.35	2.03	0.24	0.23	2.23	0.47	0.17	0.33	1.23	0.29	0.53	1.05	0.22	0.28	0.42	. . \$
,	44459983	H07920	AA070226	N66580	AA447522	AA772803	AA459945	AA598538	AA452125	AA778206	AA425319	R15779	N52350	AA974495	R51064	AA442095	W46577	AA463643	N63153	AA812996	AA181646	AA398262	R42490	AA047338	AA885471	AA702785	N72150	W57712	N53534	AA448283	AA704255	R45672	R43452	AA416782	AA885311
	796448	45578	530814	278809	782575	969568	796408	898162	786550	448379	773276	53371	284479	1372140	37283	774751	324122	811845	285261	1376853	613070	726699	31056	509495	1466418	448014	291129	340898	284220	782847	450574	35503	32381	731337	1461664
	1324																				1344														1358

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Celi-Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

-	-	-	-	- -	-	٠.		-	-	-	-	Ψ-	₩-	-	-	-	_	-	-		-	-	-	-	-	-			-	-	-	-	-	-	-		
21.07	21.75	0.77	55.	2.80	3 6	87.7	1.07	0.42	0.85	0.53	1.82	2.07	0.74	0.96	1.52	0.97	1.05	1.44	1.23	0.82	1,00	1.27	1.72	0.71	0.48	2.23	2.69	20.54	19.28	19.52	21.03	20.18	18.26	0.41	2.00		
0.49	1.49	22.43	4. 4. 4. 4.	2.1.20	20.14	19.87	22.05	21.87	20.92	22.36	21.01	1.68	21.75	22.04	20.70	21.42	21.62	20.2	20.67	20.64	21.29	20.21	19.58	20.96	21.62	18.03	18.23	1.28	1.87	1.27	0.44	0.14	1.00	20.91	18.62		
0 0	80 +	3 3	0.1	cs.1	4.14	1.92	0.90	1.66	2 14	; c	1.05	20.12	1.35	22.0	133	<u> </u>	80:- 89:0	3 6	. c.	1.2.1	c	0.0		60°		235	. F.	0.67	1.30	1 45	0.76	1 82	1 0 1 0 1 0	7.70 9.66	9 6	10:-	
		 	8.07	8.03	8.03	8.02	8.01	1 08	7.1	/s:/	7.90	7.30	7.30	7 .93	7.92	00.7	28.7	7.77	7.74	7.71	7.63	7.62	09.7	7.56	7.50	66.7	, ', , , ,	7.33	7 . 10) i	1 1	- t	0.7	7.35	8 6	(s.)	40
•	17.90	2.94	1.93	3.62	4.55	0.40		. co.	4.52	0.25	1.55	5.16	4.12	11.10	1.17	8.58	5.83	1.78	4.12	6.90	5.09	0.08	11.32	1.07	4.20	6.05	0.46	0.79	3.30	0.41	6.24	0.10	26.15	0.07	3.92	3.50	
	2.20	0.36	0.24	0.45	0.57	900	0.00	0.35	0.57	0.03	0.19	0.65	0.52	4.40	0.15	1.09	0.74	0.23	0.53	0.90	0.67	0.01	1.49	0.14	0.56	0.80	90.0	0.10	0. 44.	0.05	9. 4	0.0	3.54	0.01	0.54	0.48	
	AA452095	AA629020	AA401380	W60795	A A 46.4204	100#0147	AA449823	AA430202	H29207	R60713	R44404	N67891	H75459	N51343	AA488604	AA609749	N49186	AA426092	N47952	AA609005	AA451890	T54342	AA405532	AA447476	N62817	W85931	AA088231	N52589	AA481397	R11236	R52082	N55266	AA173189	AA668178	AA453585	AA187982	
	786504	743992	742702	3448R4		015585	788629	781404	49836	42009	34641	286661	230591	283158	843058	1031919	280213	757257	280266	1030781	786669	69272	772408	784272	278516	416158	511117	283874	746321	129392	40296	245768	595695	852829	795199	626551	
	1359	1360	138	2 5	205	1363	1364	1365	1366	1367	1368	1369	1370	1371	1372	1373	1374	1375	1376	1377	1378	1379	1380	1381	1382	1383	1384	1385	1386	1387	1388	1389	1390	1391	1392	1393	

- -	- •	,	-	-	-	-	-	-	~-	-	-	-	~	-	-	-	-	-	•		- •	- ,	-	-	-	-	-	-	-	-	-	-		-	•	•	
1.25	4 .	1.76	1.29	0.69	0.54	1.40	0.81	1.79	0.08	2.15	1.05	1.08	1.15	0.95	1.36	1.98	2.14		19.03	19.02	os:-	1.82	0.84	0.76	0.99	1.81	0.59	1.70	0.89	0.93	0.91	1.36	0.55	28	0.52		
18.94	19.21	18.29	19.38	19.82	20.04	19.41	45.95	16.61	0.29	2.76	19.69	18.84	18.55	90.91	26. 85	28.0		7.00 7.00	00.00	0.20	17.91	17.72	18.47	18.92	18.61	17.58	19.11	17.15	18.38	18 44	7 6	1 2 2	40.47	it o	17.09	T.6.	
1.72	1.18	1.76	1.12	1 22	600	26.0	5 5) 90 c	Se: 20	20.33	2 6		5 4		0.90	0.02	18.11	16.68	1.05	1.55	4.	1.16	1.39	101	1.06	1 26 1 26	S) t	<u> </u>	<u>,</u> ;	. .	51.1	0.20	0.39	0.67	0.76	
7.31	7.28	7.27	 	07.7	6. 1	7.7	ct./	7.14	21.7	7.12	۲۲.7	80.7	70.7	40.7	7.00	6.97	6.97	6.94	6.93	6.93	6.90	6.90	06)) (S 6	60.0	88.9	6.87	6.83	6.83	6.81	6.81	6.80	6.80	6.80	6.80	4
1.40	3.14	7 7	77.7	1.44	8.29	7.41	0.63	4.11	1.86	1.32	1.50	1.74	1.59	0.70	6.35	2.53	13.98	3.06	6.77	43.23	2.38	3.11	- 0 - 0	0.00	0.66	4.70	1.65	5.14	5.80	1.93	4.90	7.07	2.00	2.92	4.43	5.43	
91.0	27	54.0	rs.0	0.20	1.14	1.03	60:0	0.58	0.26	0.19	0.21	0.25	0.22	0.10	0.91	0.36	2.01	0. 44.	0.98	6.24	0.35	0.00		0.56	0.10	0.68	0.24	0.75	0.85	0.28	0.72	. 2	0.29	0.43	0.65	0.80	
A A 490240	00100000	AA620760	N57475	R59990	N91797	W88786	AA733105	AA131240	N64814.	W74802	W84778	AA478965	R51865	N63436	AA412046	AA453795	AA459980	AA191424	N51069	N67864	100,001	7,445/0/3	AA432144	N50733	R51872	AA156022	R40037	AA435988	H72107	H17516	AA621750	AA129217	N89783	AA412477	N92901	AA454170	
023772	711679	1049293	277322							345081															39266	590120	27508	730742	213634	50618	1030854	564847	302073	730146	307660	795334	
7	1394	1395	1396	1397	1398	1399	1400	1401	1402	1403	1404	1405	1406	1407	1408	1409	1410	144		2141	1413	1414	1415	1416	1417	1418	1419	1420	1421	1422	1423	1424	1425	1426	1427	1428	

		٠,	-	-		-	~	-	-	-	-	-	-	•	-	-	₩.	•	Ψ-	•	-	-	-	-	-	-	-	•	-	-	-	-	•	~~		
2.69	4.24	47.	1.09	1.19	0.74	0.73	0.63	0.67	17.37	2.35	0.83	0.76	0.38	. .	2.84	17.84	16.28	1.02	0.11	0.97	1.68	0.34	0.30	0.76	1.50	14.90	0.87	16.61	0.84	1.43	1.17	2.93	44.	2.84		
16.56	9:5	18.08	17.97	18.29	18.42	18.56	18 92	18 03	S 5 C	16.50	18.64	18.31	18.25	16.69	15.70	0.22	001	17.83	18 57	47.20	16.06	18.17	17.90	18.01	15.1	2.64	17.38	2.07	18.14	17.01	48.	14.61	17 14	14.87		
1.09	1.00	0.97	1.20	0.76	1 07		0.92	20.0	Le:0	8).	÷	- G.O.	4 27	78.	<u>;</u>	<u> </u>	1.72		0.63 0.63	5.5	- + 6 6	 		 		5 6	1.01	28.0	, t	5 5	1.01	}	60	V.07	?	
6.78	6.77	6.77	6.75	6 7	2.5	0.74	6.73	6.72	6.70	6.66	6.66	6.66	6.63 6.83	6.63 6.03	6.62	6.60	6.59	6.58	6.56	6.56	6.55	6.55	6.54	6.53	6.53	6.52	5.52		F.0 F.0	6.50	φ. τ.	0.40	6.48	6.48	0.41	42
1.64	0.07	0.71	· ·	8C.7	4.72	5.68	2.58	0.11	3.42	51.77	5.28	12.09	5.41	3.47	1.22	1.82	19.16	0.07	2.30	09:0	3.63	4.51	8.94	1.36	3.15	3.12		4.15	5.61	6.21	1.84	2.83	<u>.</u> 2	9.0	9.80	
0.24	0.01		2	0.38	0.70	0.84	0.38	0.02	0.51	7.77	0.79	1.81	0.81	0.52	0.18	0.27	2.91	0.01	0.35	0.09	0.55	0.69	1.37	0.21	0.48	0.48	0.28	0.64	0.86	96.0	0.28	0.44	0.16	0.01	43.	
04401840	TK2820	135050	AA166703	AA142923	N30562	AA461507	AA844998	AA485427	AA827369	AA46451	AA040332	N32095	AA454080	N58276	R37079	N47075	AA608856	AA677522	AA401695	AA454082	W58291	AA621363	AA456148	R45632	W78168	AA868038	W96273	W49619	AA465168	AA878739	AA708686	AA625990	R51261	AA458487	N63286	
190572	409314	70000	593652	505227	257167	795829	1412398	811046	1422424	781046	376040	260216	788234	247862	27333	280308	1048586	460398	727275	788240	341061	1055144	796369	35620	346897	1461161	361656	325182	815051	1417935	384939	745493	38528	809611	290162	
Š	1429	55 55 55	1431	1432	1433	1434	1435	1436	1437	1438	1439	44	144	1442	1443	<u>4</u>	1445	1446	1447	1448	1449	1450	1451	1452	1453	454	1455	1456	1457	1458	1459	1460	1461	1462	1463	

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line Genes upregulated 3 fold or higher in at least 2 of 3 or 10 fold or higher in at least 1 of 3 cancer cell lines.

1465 782838 AA448270 1466 731040 AA421271 1468 73524 AA453893 1469 487383 AA046705 1470 823578 AA406068 1471 743025 AA406068 1472 37404 R51100 1473 731198 AA417355 1474 303023 N91566 1475 786154 AA448855 1476 361567 AA016980 1477 727137 AA388757 1478 665033 AA194646 1479 461420 AA704943 1480 591540 AA168735 1481 731050 AA704943 1482 162734 H27545 1483 289903 N64597 1484 135900 R33609 1485 609530 AA180163 1487 277631 N45983 1488 1569463 AA973927 1489 255286	AA448270 AA421271 W60288 AA453993 AA497040 AA497040 AA417355 N91566 AA417355 AA417353	1.12 0.30 0.57 0.85 1.36 0.25 0.21 0.61 0.61 0.61 0.02	7.19 1.88 3.64 5.43 12.45 1.60 1.45 1.33 3.82 2.98 1.04 7.16	6.39 6.35 6.35 6.29 6.29 6.20 6.20 6.19	0.49 1.27 1.27 0.85 1.15 1.15 1.16 0.99 1.81 0.84	0.43 15.24 16.26 17.18 16.71 16.91 16.93 16.63 16.75 16.75 16.75 17.00 14.44 1.89	18.24 2.08 1.53 1.03 1.28 1.79 1.79 0.92 0.78 0.78 1.26 17.69	
731040 342039 795254 487383 823578 743025 37404 731198 303023 786154 361567 727137 665033 461420 591540 731050 162734 289903 135900 609530 742928 742928	21271 21271 0288 53993 46705 97040 06069 1100 17355 1566 48855 16980 98757 94646 04943	0.30 0.57 0.85 1.96 0.25 0.23 0.61 0.61 1.15 0.02	1.88 3.64 5.43 12.45 1.60 1.45 1.33 3.82 2.98 1.04 7.16	6.35 6.35 6.35 6.29 6.29 6.27 6.22 6.20 6.19	1.80 0.85 1.25 1.15 0.65 1.16 0.99 1.81 0.84 1.87	15.24 16.26 17.18 16.71 16.91 15.30 16.03 16.03 16.75 17.00 14.44 1.89	2.08 1.53 1.03 1.08 1.28 1.79 1.79 0.94 0.78 2.25 1.26	
342039 795254 487383 823578 743025 37404 731198 303023 786154 361587 727137 665033 461420 591540 731050 162734 289903 135900 609530 782668 277631 1569463 255286 461965	0288 53993 46705 97040 06069 1100 17355 1566 16980 98757 94646 04943	0.57 0.85 1.96 0.25 0.23 0.21 0.61 0.48 0.17 1.15	3.64 5.43 12.45 1.60 7.26 1.45 1.33 3.82 2.98 1.04 7.16	6.35 6.35 6.34 6.22 6.22 6.20 6.19	1.27 0.85 1.15 1.15 0.65 1.76 1.18 1.16 0.99 1.81 0.84 1.87	16.26 17.18 16.71 16.50 16.91 16.03 16.03 16.75 17.00 14.44 1.89	1.53 1.08 1.08 1.28 1.79 1.79 0.92 0.92 0.78 2.25 1.26	
795254 487383 823578 743025 37404 731198 303023 786154 361587 727137 665033 461420 591540 731050 162734 289903 135900 609530 782668 277631 1569463 255286 461965	53993 46705 97040 06069 1100 17355 1566 16980 98757 94646 04943	0.85 0.25 0.23 0.23 0.21 0.61 1.15 0.02	5.43 12.45 1.60 7.26 1.33 3.82 2.98 1.04 7.16 0.70	6.35 6.34 6.29 6.29 6.27 6.22 6.20 6.19	0.85 1.25 1.15 0.65 1.76 1.18 1.16 0.99 1.81 0.84 1.87	17.18 16.71 16.50 16.91 15.30 16.63 16.63 16.75 17.00 14.44 1.89	1.03 1.28 1.31 1.79 1.60 0.94 0.78 0.78 1.26 17.69	
487383 823578 743025 37404 731198 303023 786154 361587 727137 665033 461420 591540 731050 162734 289903 135900 609530 782668 277631 1569463 255286 461965	46705 97040 06069 11355 17355 1566 48855 16980 98757 94646 04943	1.96 0.25 0.23 0.21 0.61 1.15 0.01 0.02	12.45 1.60 7.26 1.45 1.33 3.82 2.98 1.04 7.16	6.34 6.29 6.29 6.27 6.22 6.22 6.19 6.19	1.25 1.15 0.65 1.76 1.16 0.99 1.81 0.84 1.87	16.71 16.50 16.91 15.30 16.63 16.63 16.75 17.00 14.44 1.89	1.08 1.28 1.31 1.79 1.60 0.94 1.49 0.78 1.26 17.69	
823578 743025 37404 731198 303023 786154 361587 727137 665033 461420 591540 731050 162734 289903 135900 609530 782668 277631 1569463 255286 461965	97040 06069 1100 17355 1566 48855 16980 98757 94646 04943	0.25 0.23 0.21 0.61 0.47 1.15 0.02	1.60 7.26 1.45 1.33 3.82 2.98 1.04 7.16	6.39 6.29 6.24 6.24 6.22 6.19 6.19	1.15 0.65 1.76 1.18 0.99 1.81 0.84 1.87	16.50 16.91 15.30 16.03 16.75 12.34 17.00 14.44 1.89	1.28 1.31 1.79 0.92 0.94 14.49 1.26 17.69	
743025 37404 731198 303023 786154 361567 727137 665033 461420 591540 731050 162734 289903 135900 609530 782668 277631 1569463 255286 461965	06069 117355 1566 48855 16980 98757 94646 04943	1.15 0.23 0.21 0.61 0.17 1.15 0.02	7.26 1.45 1.33 3.82 2.98 1.04 7.16 0.70	6.29 6.28 6.24 6.22 6.20 6.19	0.65 1.76 1.18 1.16 0.99 1.81 0.84 1.87	16.91 15.30 16.03 16.63 16.75 17.00 14.44 1.89	1.31 1.79 1.60 0.92 0.94 14.49 0.78 2.25 1.26	
37404 731198 303023 786154 361567 727137 665033 461420 591540 731050 162734 289903 135900 609530 782668 277631 1569463 255286 461965	1100 17355 1566 16980 16980 98757 94646 04943	0.23 0.21 0.61 0.48 0.17 1.15 0.02	1.45 1.33 3.82 2.98 1.04 7.16 0.70	6.28 6.27 6.22 6.22 6.19 6.17	1.76 1.18 1.16 0.99 1.81 0.84 1.87	15.30 16.03 16.63 16.75 2.34 17.00 14.44 1.89	1.79 1.60 0.92 0.94 14.49 2.25 1.26 17.69	
731198 303023 786154 361587 727137 665033 461420 591540 731050 162734 289903 135900 609530 782668 277631 1569463 255286 461965 742928	17355 1566 48855 16980 98757 94646 04943 58735	0.21 0.61 0.48 0.17 1.15 0.02	1.33 3.82 2.98 1.04 7.16 0.70	6.27 6.22 6.22 6.20 6.19 6.19	1.18 1.16 0.99 1.81 0.84 1.87 15.37	16.03 16.63 16.75 2.34 17.00 14.44 1.89	1.60 0.92 0.94 14.49 0.78 1.26 17.69	
303023 786154 361587 727137 665033 461420 591540 731050 162734 289903 135900 609530 782668 277631 1569463 255286 461965 742928	1566 48855 16980 98757 94646 04943 58735	0.61 0.48 0.17 1.15 0.11 0.02	3.82 2.98 1.04 7.16 0.70	6.24 6.22 6.21 6.20 6.19	1.16 0.99 1.81 0.84 1.87 15.37	16.63 16.75 2.34 17.00 14.44 1.89 0.41	0.92 0.94 14.49 0.78 2.25 1.26 17.69	
786154 361587 727137 665033 461420 591540 731050 162734 289903 135900 609530 782668 277631 1569463 255286 461965	48855 16980 98757 94646 04943 58735	0.48 1.15 0.11 0.02	2.98 1.04 7.16 0.70 2.50	6.22 6.21 6.19 6.19	0.99 1.81 0.84 1.87 15.37	16.75 2.34 17.00 14.44 1.89 0.41	0.94 14.49 0.78 2.25 1.26 17.69	ه سد سه سه سه سه
361587 727137 665033 461420 591540 731050 162734 289903 135900 609530 782668 277631 1569463 255286 461965	16980 98757 94646 04943 58735	0.17 0.11 0.02	1.04 7.16 0.70 2.50	6.20 6.20 6.19 6.17	1.81 0.84 1.87 15.37	2.34 17.00 14.44 1.89 0.41	14.49 0.78 2.25 1.26 17.69	
727137 665033 461420 591540 731050 162734 289903 135900 609530 782668 277631 1569463 255286 461965	98757 94646 04943 58735	0.11	7.16 0.70 2.50	6.20 6.19 6.17	0.84 1.87 15.37 0.41	17.00 14.44 1.89 0.41	0.78 2.25 1.26 17.69	** ** ** ** *
665033 461420 591540 731050 162734 289903 135900 609530 782668 277631 1569463 255266 461965	94646 04943 58735	0.11	0.70	6.19	1.87	14.44 1.89 0.41	2.25 1.26 17.69 1.04	4- 4- 4- 4
461420 591540 731050 162734 289903 135900 609530 782668 277631 1569463 255286 461965 742928	04943 58735	0.02	2.50	6.17	15.37	1.89	1.26 17.69 1.04	 ,
591540 731050 162734 289903 135900 609530 782668 277631 1569463 255286 461965 742928	58735	0.02		* (0.41	0.41	17.69	 •
731050 162734 289903 135900 609530 782668 277631 1569463 255286 461965 742928		000	0.15	6.17			- \$	•
162734 289903 135900 609530 782668 277631 1569463 255286 461965 742928	AA421276	0.00	6.12	6.15	1.25	16.18		-
289903 135900 609530 / 782668 / 277631 1569463 / 255286 461965 / 742928 / 897641 /	H27545	0.18	1.11	6.15	15.27	1.1	2.06	-
135900 609530 782668 277631 1569463 255286 461965 742928	N64597	0.81	4.98	6.14	1.14	15.91	1.39	.
609530 / 782688 / 277631 1569463 / 255286 461965 / 742928 / 897641 / 7	R33609	1.42	8.74	6.14	1.63	2.30	14.49	-
782668 / 277631 1569463 / 255286 461965 / 742928 / 897641 / 7	AA180163	0.47	2.86	6.13	1.01	15.63	1.75	-
277631 1569463 255286 461965 742928	AA447574	0.34	2.08	6.11	1.81	15.22	1.30	~
1569463 , 255286 461965 , 742928 , 897641 , ,	N45983	0.50	3.06	6.11	1.07	16.26	1.00	~ -
255286 461965 742928 897641	AA973927	1.82	11.09	6.09	13.75	1.73	2.80	-
461965 / 742928 / 897641 /	N23882	0.14	0.87	6.08	0.71	17.19	0.35	₹-
742928	AA779972	0.40	2.40	6.07	0.69	16.27	1,25	
897641	AA405181	0.48	2.91	6.07	1.59	15.36	1.25	- -
֡֜֜֜֜֜֜֜֜֜֜֜֜֜֜֜֜֜֜֜֜֜֜֜֜֜֜֜֜֜֜֜֜֜֜֜֜	4A496792	0.25	1.54	6.07	1.05	16.43	0.72	
1469857	AA865150	0.65	3.94	6.05	1.37	15.22	1.56	-
770989	VA427404	0.03	0.16	6.05	0.85	2.37	14.92	-
214624	H71242	0.52	3.17	6.05	2.52	14.36	1.26	-
280214	N49181	1.02	6.12	6.01	1.92	13.20	2.90	-
46715	H10011	0.27	1.59	6.00	2.75	1.81	13.42	-
324345	W47585	0.01	90.0	5.98	1.76	15.18	1 .00	₩-
				43				

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Lines Genes upregulated 3 fold or higher in at least 1 of 3 cancer cell lines.

-	-	-	-	-	-	-	-	-	-	-	-	Ψ-	-	-	-	+ -	-	-	-	-	-	-	-	-	-	-	+-	-	-	-	-	-	-	•		
0.86	0.75	17.23	0.34	1.12	2.48	0.74	16.74	1.90	0.40	2.85	0.28	13.10	1.22	0.88	0.67	1.27	0.87	16.0	2.69		14.81	0.56	96.0	22.0	0.18	0.07	1.79	13.64	1.08	1.31	2.88	2.58	14.84	2 .		
15.89	15.78	0.01	16.22	15.41	14.53	15.90	0.54	15 OS	0.50	13.14	14.95	3.00	15.23	15.93	15.63	15.07	2,49	£.43		13.52	9.66	45.55 F. B. 1	10.01	. 4 4 5 7	16.13 16.13	15.29	14.20	2.35	15.05	14.61	13.08	11 46	6 6	14 64	Š:	
1.18	1.32	08.0	1.28	1 27	72.0		1.10	2 6	0.72	1 67	7.42			0 - c) (57.1	80.4L	88.0	1.07	1.36	1.77	9	1.37	0.26	1.68	† †	 	. c			5.6	2.80 0.00	5 4		
5.98	, r o, r) u	ე.ფე	0. e0		 			5.90	5.8g	0 0	O 0	0.00	5.07 10.07	9.67	5.87	5.86	5.81	5.81	5.76	5.76	5.75	5.74	5.72	5.72	5.72	5.70	5.70	D 10	9.6	5.66	5.65	5.63	5.61	5.59	4
20.05	89.3	4.00	3.98	2.17	2.60	1.49	4.69	12.14	1.35	2.18	2.11	0.21	37.89	2.08	6.90	6.10	1.46	2.46	2.13	2.71	1.60	4.87	5.93	6.22	24.55	0.32	0.76	3.16	27.34	6.19	2.93	2.33	3.19	90.0	2.90	
Š	\$:	0.79	0.67	0.36	0. 4	0.25	0.79	2.06	0.23	0.37	0.36	0.0 40.0	6.44	0.35	1.18	<u>5</u>	0.25	0.42	0.37	0.47	0.28	0.85	1.03	1.09	4.29	90.0	0.13	0.55	4.80	1.09	0.52	0.41	0.57	0.01	0.52	
	AA457528	AA401409	N21576	AA479928	AA453256	AA775405	AA448825	T72235	AA628132	H24329	AA418004	H09616	AA774678	N70791	AA130955	N95476	AA447724	H43707	AA620927	N48901	N49619	R40780	W88999	N23652	AA443958	AA133469	N33366	N50729	AA485653	AA171426	H96605	AA463225	W95118	N33778	W58209	
	838500	743184	266146	772938	795339	878097	786107	85840	1055727	51749	767461	46183	856388	298162	586923	310054	813807	184256	1055636	279583	277785	28106	417461	254564	757190	586796	270681	283702	840404	594806	251517	797048	415191	272951	341095	
	1499	1500	1501	1502	1503	1504	1505	1506	1507	1508	1509	1510	1511	1512	1513	1514	1515	1516	1517	1518	1519	1520	1524	1522	1523	1524	1525	1526	1527	1528	1529	1530	1531	1532	1533	

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line Cennes, upregulated 3 fold or higher in at least 2-10: Prostate Cancer Cell Lines vs. 2 of 3 or 10 fold or higher in at least 1 of 3 cancer cell lines.

-	•	 -	-	-	-	-	-	-	₩.	٠-	-	•	-	-	₩-	-	-	-	₩		₩	-	₩.		-	~	-	-	-	-	-	-		- -	
1.39	2.72	1.44	0.34	0.43	2.93	0.80	10.87	1.76	0.34	0.54	14.48	0.93	2.71	14.43	0.64	0.98	0.77	1.59	0.69	13.75	10.73	0.14	0.52	.34 .34	0.22	1.99	0.11	1.81	0.95	1.12	1.40	1.52	12.24	09:0	
14.07	12.89	0.45	13.78	14.42	12.33	14.57	2.90	1.81	14.66	15.17	1.00	14.52	12.03	1.14	15.22	14.23	14.78	13.47	14.20	1.33	2.97	13.71	15.03	13.09	14.13	11.67	15.56	1.81	14.19	14.02	13.69	12.04	2.43	14.21	
1.31	1.17	14.87	2.63	1.89	1.47	1.33	2.92	13.10	1.63	0.90	1.11	1.11	1.82	0.93	0.59	1.22	0.88	1.35	1.46	1.27	2.64	2.49	0.78	1.86	1.92	2.61	0.59	12.59	2.5	2.7	1.07	2.60	1.48	1.30	
5.59	5.59	5.59	5.58	5.58	5.58	5.57	5.56	5.56	5.55	5.54	5.53	5.52	5.52	5.50	5.48	5.48	5.48	5.47	5.45	5.45	5.44	5.44	5.44	5.43	5.42	5.42	5.42	5.41	5.39	5.39	5.39	5.39	5.39	5.37	45
1.02	6.24	40,1	0.37	2.21	2.46	2.93	7.23	0.42	0.65	5.82	0.06	3.10	0.16	6.39	0.57	3.15	4.04	1.09	4.31	6.45	12.51	0.40	0.22	3.92	0.25	0.33	0.28	0.03	4.66	3.15	3.29	2.17	13.33	1.45	
C 87	1.12	0.19	0.07	0.40	0.44	0.53	1.30	0.08	0.12	1.05	0.01	0.56	0.03	1.16	0.10	0.57	0.74	0.20	0.79	1.18	2.30	0.07	8	0.72	0.05	0.06	0.05	0.01	0.86	0.58	0.61	0.40	2.48	0.27	
A A E O B O 7 E	AA620463	N51002	H06249	AA404356	AA186613	M95414	AA598601	H12338	R28660	N50432	H51549	AA401472	AA401470	AA703147	W73523	AA600284	T96913	H09541	W95409	W92011	H56028	AA447547	AA620472	N90514	AA009763	AA459702	R36989	AA460328	N48078	N47425	AA176607	AA205803	AA496544	H89713	
88000	051102	281243	44287	758370	625616	358212	898218	148469	133864	280633	179753	742589	742590	434845	344505	1031552	121409	45922	358200	415264	203544	782611	951101	306121	429773	795585	27098	795788	281756	280672	609052	645332	755904	240273	
7697	1534	1536	1637	153	1530	2 2 2	<u> </u>	5 2	15.43	154	1545	1546	1547	1548	45.40	בר בר ה ה	1884	1551	1553	1554	1888	1556	1557	1558	1559	1560	1561	1562	1563	1564	1565	1566	1567	1568	

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Lines Genes upregulated 3 fold or higher in at least 1 of 3 cancer cell lines.

-	-	-	-	-		-	-	-	-	-	-	-	-	-	-	-	-	-	• 🕶	- •	-	Ψ	₩.	-	-		-	-	-	-	-	-	-	• •		-		
2.86	1.50	1.31	1.13	0.97	0.61	15.12	0.38	1.43	0.97	0.76	13.96	2.04	0.63	0.49	1.96	0.83	0.47	138	3 6	0.0 0.0	1.05	1.33	14.83	 \$	2.39	40.	0.84	0.51	2.66	2.91	0.09	1.28	0.40	9 9	04.1	1.80		
10.35	13.41	1.23	13.90	13.96	14.31	81.0	44.89	43.87		14.65	1.66	4.8	13.86	14.51	1.84	13.61		- c	0.22	13.86	13.60	13.15	0.41	13.78	12.98	12.77	12.97	15.02	12.55	11.43	14.87	4.07	17:71	0.93	12.97	12.99		
2.88	1.17	13.49	2 0	6.0	8.7) · · ·	0.00	9.0	0.62	o (2)	9 6		4 27		3 5	ţ ;	D4.F	14.20	14.23	1.12	1.10	1.26	0.49	0.59	33	88	50. t	5 .	2 7	- 4. 2. 4.	0	. c.	40.	14.21	1.16	0.73		
5.36	A 36	0.00	٠ ن ن	φ 4	5.34	5.32	5.32	5.31	5.31	5.31	5.31	5.31	5.30	5.29 0.00	5.29	5.28	5.28	5.28	5.28	5.27	5.25	5.25	200	17.0 10.4	0.75 0.75 0.75	6.6	5.23	5.23	27.5	5.21	5.20	5.20	5.19	5.18	5.18	5.17	97	‡
3	3 6	2.10	1.33	4.00	0.82	2.21	6.86	4.78	9.92	20.08	4. 48.	0.18	1.87	11.30	2.08	5.76	2.96	3.67	3.47	5.25	14.4	; c	76.7	51.0	9.37	5.67	2.23	2.30	1.16	0.94	2.50	0.56	2.70	3.62	13.39	5.75) ;	
Č	٠.00	0.39	0.25	0.75	0.15	0.42	1.29	0.90	1.87	3.78	0.82	0.03	0.35	2.14	0.39	1.09	0.56	0.70	0.66	9 5	9:-	9. (0.56	0.05	1.79	1.08	0.43	0.44	0.22	0.18	0.48	0.11	0.52	0.70	2 50) , ,	-	
	AA504617	R08292	AA699895	AA211565	N29796	AA406226	R09729	W74216	AA098867	N70038	AA485714	H96643	AA626275	AA777024	R59618	N66085	AA191518	R01739	A A 406140	AA460148	AA16/338	AA609368	AA677200	N47208	AA015819	AA404269	AA682439	AA195668	AA161466	AA670359	AA705184	N20602	N50787	H10242	343611	H2/980	AA039713	
	825583	127242	462468	562158	270766	742887	128058	346359	489535	297878	84044	251591	745072	378242	42120	278386	627306	124148	757000	777/0/	595623	743452	454698	280412	360644	758309	462963	665154	592497	878417	462729	264157	283890	54044	11016	162533	376435	
	1569	1570	1571	1572	1573	1574	1575	1576	1577	1578	1579	1580	1581	1582	1583	1584	1585	2 4	200	1587	1588	1589	1590	1591	1592	1593	1594	1595	1596	1597	1598	1599	1500	3 6	5	1602	1603	

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in at least 2 of 3 or 10 fold or higher in at least 1 of 3 cancer cell lines.

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1.80	96.0	1.16	0:20	0.82	1.02	1.28	2.19	1.07	0.68	2.21	1.10	2.91	2.00	0.98	1.23	0.52	1.01	1.21	0.75	0.88	1.08	1.62	1.78	1.09	0.93	1.00	1.89	0.93	1.15	1.33	1.64	0.98	0.73	11.45	
12.54	13.61	12.96	13.19	14.23	13.32	12.49	2.33	13.03	13.91	2.45	13.20	1.75	2.00	13.11	13.43	13.26	13.06	12.47	13.28	11.90	13.49	10.93	11.65	11.42	12.41	1.00	11.91	12.76	13.08	0.13	12.45	12.80	13.67	2.40	
1.17	0.92	1.37	1.80	0.43	1.12	1.67	10.92	1.32	0.80	10.73	1.08	10.73	11.35	1.25	0.66	1.51	1.21	1.55	1.14	2.38	0.60	2.60	1.69	2.55	1.70	13.03	1.21	1.31	0.73	13.44	0.78	1.05	0.43	96.0	
5.17	5.17	5.16	5.16	5.16	5.15	5,15	5.14	5.14	5.13	5.13	5.13	5.13	5.12	5.11	5.10	5.10	5.10	5.07	5.06	5.06	5.06	5.05	5.04	5.02	5.01	5.01	5.00	5.00	4.99	4.97	4.96	4.95	4.94	4.94	47
1.38	12.87	3.90	3.54	2.63	3.04	2.68	2.14	3.22	3.64	10.18	5.71	1.36	12.23	2.19	5.54	0.87	4.24	2.18	12.39	1.78	4.31	2.25	14.29	4.11	2.63	0.05	2.76	0.98	5.69	7.99	2.23	3.75	1.17	27.18	
0.27	2.49	0.75	0.69	0.51	0.59	0.52	0.42	0.63	0.71	1.98	1.1	0.27	2.39	0.43	1.09	0.17	0.83	0.43	2.45	0.35	0.85	0.45	2.84	0.82	0.52	0.01	0.55	0.20	4.1	1.61	0.45	0.76	0.24	5.50	
07007844	A A 405809	AA609594	R16259	AA866029	AA417915	W58191	H79035	AA428659	AA608974	R92281	N40976	AA707615	AA101777	N48590	AA486084	N62696	AA400389	N49109	N91175	AA460963	N45223	AA455096	N66178	N30597	N26993	AA018683	N67619	T98056	N67570	AA432075	W72920	AA004946	AA455904	H73947	
1475972	742898	1031736	53081	1470278	767414	341065	233712	781447	1030764	196189	279827	1292121	490649	279278	840786	288936	742607	279806	301842	796123	283122	812293	278572	257746	257779	362795	290680	119289	285680	784130	345118	428298	813242	232789	
1604	100.4	1608	1607	1608	909	25.4	161	1612	1613	1614	1615	1616	1617	1618	1619	1620	1621	1622	1623	1624	1625	1626	1627	1628	1629	1630	1631	1632	1633	1634	1635	1636	1637	1638	

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Lines Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

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0.36	1.52	0.20	1.44	2.29	0.96	2.74	2.25	0.44	t 9	0.0	0.22	o 1	0.05	12.40	0.30	4.14	0.45	11.76	1.55	0.81	1.55	2.25	12.57	1.25	40.7	11.69	1.78	0.13	11.72	1.48	0.74	1.92	0.71	1.48	1.68	
13.93	12.51	14.40	12.16	11.25	12.51	40.04	0.00	10.12	15.40	12.92	12.52	13.24	0.18	1.08	12.12	11.90	12.60	0.46	11.70	12.99	11.94	11.25	1.00	12.19	12.34	1.07	11.69	12.71	1.93	11.63	11.76	11.57	12.76	11.91	1.98	
0.53	0.78	0.21	1.19	1.24	; ; ,	5 6	D (0	2.39	0.82	1.14	1.98	0.99	14.47	1.09	1.56	1.60	1.59	2.38	1.34	0.79	1.09	1.07	1.00	1,10	1.16	1.76	<u>5</u>	1.64	0.83	1.36	1.96	0.97	66.0	1.07	10.78	
4.94	4.94	4 93	4.93	 		4.92	4.92	4.92	4.91	4.91	4.91	4.90	4.90	4.89	4.88	4.88	4.88	4.87	4.86	4.86	4.86	4.86	4.86	4.85	4.84	4.84	4.84	4.83	4.83	4.82	4.82	4.82	4.82	4.82	4.82	48
72	2 23	55.5	47.11	2.4/	90.1	6.40	0.81	5.51	0.95	3.35	1.12	4.63	1.16	1.19	0.19	1.31	5.56	6.82	1.82	8.97	13.01	0.65	0.05	2.06	3.08	8.56	3.08	0.36	2.05	3.23	12.92	2.35	5.41		15.82	
ć	? ? ?	80.0 0.0	2.28	0.50	2.37	1.30	0.16	1.12	0.19	0.68	0.23	0.94	0.24	0.24	9.0	0.27	4.	1.40	0.37	48.	268	0.13	2 6	0.43	0.64	17	0.64	0.08	0.42	0.67	2.68	0 49	7	7 7	5 K	2
1	R67259	AA156112	W79445	AA406083	AA454652	AA486864	AA883680	AA206865	N35489	AA167500	H11071	AA412247	AA101632	H15695	N50563	AA421054	AA010388	0000000	4446876	A A 0.44903	1812763	20175	79090	02000	T86937	D89700	N64603	AA417275	Ho3328	W85709	A B 0.45 57.4	A 4 400694	100004	AA609334	N77779	0610070
1	41558	589861	346997	743058	811899	841022	1466633	648056	272169			731404	490484	49203	280763	731393			200000			344008	166346	26699	115277	167076	2800080	731205	242037	415976	FOORBA				289666	05/+00-
	1639	1640	1641	1642	1643	1644	1645	1646	1647	1648	1649	1650	1651	1652	1652	200	100 A	200	1000	202	0 6	8001 6001	000	1001	7001	200	1004	1000	1687	1001	900	200	0/91	1671	1672	5/91

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line. Genes, upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

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1.32	1.36	11.14	11.85	10.93	0.38	0.29	1.79	0.71	1.56	0.86	0.60	1.12	1.55	0.57	1.27	0.60	0.42	0.88	1.16	1.07	0.60	12.09	0.62	1.17	0.41	11.48	4 .	1.18	1.08	10.93	0.94	12.11	1.41	1.00	
12.21	1.36	1.32	0.33	1.75	12.91	13.94	12.05	12.67	10.53	12.47	13.16	11.33	11.93	12.95	11.28	12.29	11.88	12.35	11.65	0.73	11.61	1.53	12.86	11.17	12.67	0.35	11.08	11.59	11.34	1.15	11.22	0.01	11.01	11.30	
0.92	11.66	1.89	2.18	1.67	1.06	0.10	0.47	0.94	2.16	0.91	0.47	1.77	0.70	0.64	1.57	1.23	1.82	0.87	1.26	12.24	1.83	0.40	0.54	1.66	0.88	2.12	0.90	1.05	1.39	1.72	1.65	1.67	1,32	1.45	
4.81	4.79	4.79	4.78	4.78	4.78	4.77	4.77	4.77	4.75	4.75	4.74	4.74	4.73	4.72	4.71	4.71	4.71	4.70	4.69	4.68	4.68	4.68	4.67	4.67	4.65	4.65	4.61	4.61	4.60	4.60	4.60	4.59	4.58	4.58	49
3.96	9.	25.60	6.99	260.67	2.01	21.07	0.65	5.60	0.31	3.40	3.45	4.38	6.86	1.32	2.04	5.63	0.47	3.89	3.20	2.71	0.92	5.40	5.18	3.00	5.43	9.34	0.82	5.04	2.73	4.90	9.37	103.69	1.74	1.38	
0.82	0.0	5.35	1.46	54.50	0.42	4.4	0.14	1.17	0.0	0.72	0.73	0.92	1.45	0.28	0.43	1.20	0.10	0.83	0.68	0.58	0.20	1.15	1.	0.64	1.17	2.01	0.18	1.09	0.59	1.06	2.04	22.57	0.38	0.30	
04461119	R35665	H27379	44598587	WO5026	AA005358	AA278759	AA057620	AA844447	AA609422	AA459857	AA779048	N49098	N59295	N68928	AA634482	AA452578	AA454654	AA461086	AA431975	N30096	W20462	N91811	AA195651	H05777	AA011597	AA417895	T89084	N39590	AA401436	AA877815	AA486445	AA873060	AA412485	W73022	
796199	137017	163174	808105	295483	429202	703581	377051	1390860	743536	795810	858761	279771	289794	297857	743901	788519	811907	796170	782208	256907	327480	306513	665148	43936	429681	752770	22541	277063	743193	1161564	842927	1476065	730343	345318	
1674	1678	1678	744	4578	1670	1680 C880	1681	1682	1683	1684	1685	1686	1687	1688	1689	1690	1691	1692	1693	1694	1695	1696	1697	1698	1699	1700	1701	1702	1703	1704	1705	1706	1707	1708	

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: denes tipregulated 3 fold or higher in at least 1 of 3 cancer cell lines.

-	-	-	•	- ,		•	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	₩-		-	τ-	-	-	-	-	-	-	-	-	-		
1.70	1.05	1.26	- +	S. 1	0.28	0.45	£	11.60	2.11	2.26	2.26	0.51	0.44	0.98	1.64	1.50	1.29	0.76	11.46	0.39	1.39	0.71	1.00	2.45	0.87	0.63	2.19	40.1	0.77	1.31	1.03	0.21	1.12	0.90	0.48		
11.01	11.06	0 0	11.30	11.52	12.54	12.10	11.03	0.90	26.0	10.76	10.02	12.14	12.07	11.61	10.60	11.03	10.50	11.58	9	12.03	10.20	12.07	0.61	10.04	11.55	11.78	10.29	10.92	11.30	10.60	11.12	12.55	11.25	11.45	0.39		
0.97	. t	26.	0.98	0.84	0.79	1.05	1 23	2	10.4 45.45	54.5	1.26	88.0	50.5		1.26	7 . C	, d	9 5	1.14	8. 5) 84.0	11.75) w	8 6	9 6	28.0	46.	121	25		0.49	9 8 8 5 C	0000	12.36	3	
7	00.4	4.55	4.54	4.54	4.54			4 		15.5	10.4	10.4	10.4	10.4	4.50 6.40	4. 9. (4.49	4.49	4.49	4,49	4.48	4.46	04.4 C 14.	C4.4	4.45	4	4. 4.	4. 4. 5. 6.	r r	4. 4	7.4.	74.4	4.42	4.41	7.4	4.41	20
•	3.82	2.28	2.94	3.70	9 6	5.	4.48	4.06	4.46	1.66	2.99	1.40	3.80	2.04	0.60	2.03	2.17	3.23	4.55	0.0	6.20	80. 40.	2.59	2.85	2.32	3.08	1.74	9.0	08.1	1.88	-	4.58	4.12	1.25	5.27	6.26	
	0.84	0.50	0.65		0.02	0.88	0.99	0.90	0.99	0.37	99.0	0.31	0.84	0.45	0.13	0.45	0.48	0.72	1.01	0.01	1.38	1.80	0.58	9.0	0.52	0.69	0.39	0.15	0.43	0.42	0.25	4 .	0.93	0.28	1.19	1.42	
	N90583	AA609430	H07644	100	AA454844	W92798	AA702788	AA487046	AA703117	H17022	AA486288	AA634479	AA454085	AA398247	R46700	W79525	AA666087	AA101833	N22210	N63768	W86832	AA464861	AA682549	AA777098	R98407	N51574	H11631	T58430	AA453261	N59870	AA085759	AA412499	AA427947	R39014	R22024	N31952	
	306243	743531	200	717107	809976	418400	448024	841153	434828	50781	842847	743893	788246	726663	36491	347066	859382	489549	254468	293013	416889	789383	431255	377898	200656	280431	48060	75644	795349	284803	488276	730147	773512	24895	130742	259374	,
	1709	7	2 :	1711	1712	1713	1714	1715	1716	1717	1718	1719	1720	1721	1722	1723	1724	1725	1726	1727	1728	1729	1730	1731	1732	1733	1734	1735	1736	1737	1738	1739	1740	1741	1742	1743	

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Cine. Series upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

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10.87	10.55	12.75	10.97	10.93	1.00	1.23	1.63	1.37	0.78	0.62	1.54	1.29	0.31	0.73	0.65	1.39	0.82	0.97	76.0	1.02	0.69	0.84	0.68	0.76	0.79	0.96	0:30	11.65	11.10	1.25	0.52	1.07	0.75	0.96	
0.21	1.71	0.18	0.11	0.08	11.12	10.89	1.12	10.22	10.88	11.31	10.36	10.40	11.87	11.07	11.68	10.57	10.98	10.85	10.98	10.44	10.60	10.91	11.45	11.09	10.13	10.76	10.89	0.63	0.72	10.21	10.85	10.41	0.38	10.46	
2.15	0.94	0.21	2.06	2.12	0.98	0.99	10.31	1.46	1.39	1.1	1.13	1.33	0.81	1.18	0.63	66.0	1.09	49.	0.87	1.36	1.51	2.	0.65	0.92	1.83	1.02	1.53	0.41	0.82	1.17	1.25	1.14	11.47	1.18	
4.41	4.40	4.38	4.38	4.38	4.37	4.37	4.35	4.35	4.35	4.35	4.34	4.34	4.33	4.33	4.32	4.32	4.29	4.29	4.28	4.27	4.27	4.26	4.26	4.25	4.25	4.24	4.24	4.23	4.21	4.21	4.21	4.21	4.20	4.20	51
0.21	5.48	18.96	0.38	16.86	0.68	0.83	2.03	4.05	7.94	2.37	1.03	3.50	0.73	2.12	2.84	1.27	0.65	2.39	2.36	5.35	4,38	4.69	6.00	2.81	2.56	2.74	0.10	4.66	11.90	0.86	1.98	3.22	2.79	2.71	
0.05	1.25	4.33	0.09	3.85	0.16	0.19	0.47	0.93	1.83	0.55	0.24	0.81	0.17	0.49	0.66	0.29	0.15	0.56	0.55	1.25	103	1.10	4.	0.66	0.60	0.65	0.02	1.10	2.82	0.20	0.47	0.77	99.0	0.65	
44001222	D28254	H87471	H18433	AA167269	AA620783	P00831	A 4 7 7 8 0 8 9	AA187679	AA169121	AA609135	AA705103	AA609744	AA463444	N63543	A B 278731	PA2560	R43805	4460702	AA457474	4448172	0/020/0/	N22495	44417250	A418414	AA447743	N34892	AA398267	T60168	N31564	R41804	N91382	N51583	AA431988	AA454186	
976698	124666	252515	50986	595604	1049321	201045	27070	625838	50055 504556	1031367	462598	1031907	811764	278198	1417034	30821	35410	796711	838278	782773	418384	254276	731183	767284	813639	276689	726709	81427	271672	30894	306005	280453	782217	795452	,
777	47.65	1746	1747	1748	1740	140	47.54	157	4769	784	1754	47.5	1757	4758	2 4	1780	4764	1787	1763	787	1784	447	1787	1758	1769	4770	1771	1772	1773	1774	1775	1776	1777	1778	

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithellal Cell Line Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

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0.52	1.12	1.07	0.49	0.81	12.16	0.91	0.89	88.0	0.07	(8.0 C8.0	10.41	1.75	 	35	0.82	0.76	2 0	6.0	15.0	10.18	0.53	10.22	0.23	1.26	11.28	0.28	0.76	0.91	1.03	1.25	0.72	0.56	7 7 7	6.6	F8:0	0.62	
10.99	0.35	10.39	0.51	0.02	0.15	10.52	10:01			10.02	10.33	9.7	1 10	10.78	10.05	0.0	10.58	70.32	10.47	1.18	10.08	0.02	11.66	10.38	0.27	11.02	10.39	10.21	10.27	10.00	40.15	10.15	5 6	10.42	10.16	10.43	
1.08	11.10	1.10	11.57	11 74	25.0	2.5	7	1.35	40.0	1.41	1.25	0.93	1.4.0	74.0	0.89	0.83	0.93	0.95	1.43	0.85	1.60	1.93	0.28	0.52	0.59	0.82	96.0	96.0	0.75	0 6	 	o :- 6	S :	0.96	0.80	0.80	
4.20	4.19	4.19) (9 C	. t	9 9	81.4	4.18	4.17	4.13	4.13	4.11	4.10	4.10	4.10	4.09	4.09	4.09	4.07	4.07	4.07	4.06	4.06	4.06	4.04		t 50.5	5 6	1 0	2.02	D.4.	4.01	4.00	3.98	3.96	3.95	52
57 %			6/:-	9.4.0 9.1	2.67	38.53	3.65	4.43	0.35	3.11	4.41	0.04	1.03	2.80	2.10	2.74	2.71	1.43	0.13	2.12	6.21	8.26	3.16	14.17	08 %		3.07		77.1	1.08	0.90	2.00	8.94	1.14	2.21	6.19	
ç	3	5.5.	0.42	1.55	0.64	9.21	0.87	1.06	0.08	0.75	1.07	0.01	0.25	0.68	0.51	0.67	99.0	0.35	0.03	0.52	1.53	20.	0.78	; ;	n 4	0.60	0.76	1.53	0. 44.	0.27	0.22	0.50	2.24	0.29	0.56	1.57	
	AA453465	N91003	AA412446	N47941	W95082	AA620995	AA421001	W81117	AA449686	AA455130	AA470079	T50995	AA778533	W80791	AA455358	AA707753	R05660	AA425749	AA479912	H20046	A A 4 4 7 5 6 1	AA487146	A4407140	AAC826034	AA187287	AA701933	R15891	AA443823	AA399633	H08560	AA621478	AA136071	N78903	AA194189	P37780	AA159613	
1	795168	306066	730100	280217	415145	1056172	731298	415413	785760	809863	730406	76647	1048872	347268	812222	412983	29397	773392	772916	172785	782617	044053	04120	00007/	625465	435573	53122	784064	729924	45542	1055278	490071	300024	665738	27072	593218	
	1779	1780	1781	1782	1783	1784	1785	1786	1787	1788	1789	1790	1791	1792	1793	1794	1795	1796	101	130	1 20	88.	0081	1801	1802	1803	1804	1805	1806	1807	1808	1809	1810	181		1014	2

-	-	Ŧ	•	-	Ψ-	• •	-	-	-	• •	-	-	•		- -	-	•	٠ ,	-	-	-	
1.00	0.79	6	77.0	0.68	40.05	200	0.14	0.43	78.0	5	10.13	10.44	77.0	t S	0.48	0.64	0.40	P i	0.05	0.25	0.32	<u> </u>
10.07	62.0	3	11.14	10.25	300	Č.	10.29	10.27	9 6	81.01	0.61	0.20		10.26	10.60	10.08		0.12	0.80	10.04	5 0	i i
0.76	30.04	0.50	0.46	78.0	5	0.74	1.30	000	0.00	0.68	0.78	07.0	2.5	0.42	0.25	93.0	90.0	10.66	10,29	3 8 6	6.0	0.32
3 94) i	3.94	3.94		5.0 T	3.91	4 94	5 6	3.80	3.84	2 84	5 6	3.81	3.81	3 78) i	3.76	3.72	4 70) i	3.65	3.60
ç	7.40	0.05	80 0	9	3.36	96.6	1). O	10.36	0.92	7 1	4.0	0.19	0.93	9 6	0.42	4 .06	24.47	ç	c/:	0.15	0.11
;	0.63	0.01		0.20	0.85	2.55		0.20	2.69	0.24		0.19	0.05	76.0	7	0.11	1.08	6.57	; !	0.47	9.0	0.03
	AA150401	A A 4 7 9 9 0 6	2000	AA432064	AA609056	A A 470242	278182	H08598	AA863169	DE0747	71 7002	AA460285	AA777886	10000	AA42000/	R60705	AA885470	0003000	AA623080	AA176867	AA007687	AA148735
	491690	772004	1007	784105	1031279	10.000	104555	45783	1455671	10446	42115	795736	449438		757238	41903	1466416	77.07.1	744844	609155	429424	504761
	1814	404	0	1816	1817		1818	1819	1820		1821	1822	4022	3	1824	1825	1826		1827	1828	1829	1830

Table 2-11: Prostate Cancer Cell Lines vs. Normal Prostatio Epithelial Cell Line Cenes upregulated 5 fold or more in at least 2 cancer cell lines.

Table 2-11: Prostate Cancer Cell Lines vs. Normal Prostastic Epithelial Cell Line: Genes upregulated 5 fold or more in at least 2 cancer cell lines

Order	IMAGE Clone ID	Gen Bank Accession Number	MPM1(PrEC)	Ave Cancer Cell Lines Values	Fold(AveCell line/MPM1)	Fold(MPM2/MP Fold(MPM271/M # of Cancer M1) PM1) Cell Lines	Md(MPM271/M F PM1)	old(MPM272/M PM1)	# of Cancer Cell Lines
	1		č	7. A	591.67	7.33	66.37	1701.31	က
-	757165	AA443950	0.0	- (03.4.30	8.10	680.66	14.13	က
7	592707	AA160606	0 .01	00.	406 56	9 70	570.75	10.82	ო
က	220851	H95633	0.01	1.28	196.30	. œ	526.08	7.15	ო
4	782804	AA448186	0.01	1.81	180.70	0000	304 13	22.85	က
w	1343732	AA725564	0.01	1.12	112.48	0.45.0 0.45.0	201.10	43.45	ო
9	784296	AA447079	0.01	0.66	65.30	45.84	100.00	35.70	· m
7	796569	AA460463	0.01	0.35	63.57	16.10	138.91	33.70) (°
. 00	1558655	AA976561	0.08	4.36	54.32	92.06	17.92	18.74) m
, σ	743016	AA406036	0.01	0.50	50.47	15.70	123.80	11.93) (
, c	39442	R51617	0.10	5.02	49.93	99.52	44.77	5.49	o (
÷ ‡	33076	R44048	0.80	36.21	45.35	56.36	29.19	F6.06	? (
: \$	626300	44189108	0.07	3.16	45.16	91.62	20.97	22.90	n (
2 9	060000	7703100	500	4.0	44.34	25.03	101.07	6.93	, ო
2 ;	200071	760274	5 6	030	37.18	16.25	80.79	14.51	ო
4 ;	0/1218	AA456035	5 6	0.32	36.79	12.25	87.45	10.68	ო
.	20288	H1/888	5 6	3 6	35.62	5.91	88.28	12.68	ო
5	243181	H94482			34 44	20.19	73.72	9.41	က
17	256911	N30098	10:0	0.20	33 18	5.83	82.72	11.01	ო
₩	1030613	AA608832	D.O.	0.55	2 6	20.44	39 44	28.82	က
19	487861	AA045436	0.02	0.59	32.57	t :67	9.24	76.87	ю
20	270626	N33331	0.01	0.32	31.52	6. 50 6. 50 6. 50	38.30	24.80	ო
73	291057	N72115	0.23	6.85	29.19	24.40	9.00	48 14	m
23	1455835	AA863292	0.01	0.40	29.16	9.10	30.24		» «
23	108378	1777729	0.01	0.41	29.13	7.73	36.47	£5. 5) (
2,	1391682	AA789328	0.02	0.44	28.83	5.99	65.78	14.72	? (
, ç	263014	H99813	0.63	17.34	27.31	19.07	21.59	41.29	n (
3 8	303180	N92764	0.0	0.27	27.21	13.82	36.04	31.77	ro ·
3 6	42445	DAGGR4	0.12	3.11	26.71	9.46	65.35	5.30	ო
7) i a 7 a	20000	!		•				

Table 2-11: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 5 fold or more in at least 2 cancer cell lines.

က	က	က	ო	ო	ო	က	က	က	ო	က	က	က	က	က	m	rs (m	ເ ວ (თ ((A)	က ·	ლ -	က	က	с	က	က ⁽	m	ო	ო	က	က	က	က	
36.33	13.87	25.64	14.11	12.48	26.60	10.88	32.50	9.98	26.43	11.55	49.72	24.62	22.42	9.30	40.27	20.16	14.13	10.73	16.88	¥.94	31.84	32.94	5.91	33.48	9.40	26.82	12.25	6.49	5.36	15.00	22.27	5.13	26.44	5.25	
35.03	30.16	13.26	58.23	52.20	13.61	16.26	10.09	17.55	34.50	16.96	7.89	18.45	33.42	30.57	11.18	24.35	28.90	33.82	8.91	11.78	10.67	8.92	36.98	5.09	30.25	15.89	14.07	33.79	5.66	24.61	16.46	32.86	6.51	31.74	
7.36	33.86	38.89	5.18	12.08	34.18	46.38	30.26	44.37	8.11	38.20	5.07	18.18	5.36	20.94	7.04	10.20	10.45	7.52	25.95	5.64	8.16	7.84	6.72	10.47	9.38	5.81	21.48	6.61	33.97	5.14	5.11	5.64	9.65	5.44	
26.24	25.96	25.93	25.84	25.59	24 80	24.51	24.29	23.97	23.01	22.24	20.90	20.42	20.40	20.27	19.50	18.24	17.83	17.36	17.25	17.15	16.89	16.56	16.54	16.35	16.34	16.17	15.94	15.63	15.00	14.92	14.61	14.54	14.20	14.14	2
0.16	121	12.53	22.0	0.47	5 5	S); 0	9.30	0.16	9 0	3.52	0.21	0.25	0.20	2.46	0.19	0.31	2.95	0.21	33.78	0.63	0.17	0.17	0.21	0.16	60:0	11.31	0.32	0.29	3.80	0.35	0.35	0.14	0.14	0.50	
č	. v	9 6 8 8	5 6	5 6	5 6	5 6		5 6	5 6		2 5	50.0	6	0.12	0.0	0.02	0.17	0.0	1.96	9	0.01	0.01	000	00	0.0	0.70	0.02	0.02	0.25	0 0	0.02	0.0	0.0	9.	
	AA416903	H09910	H02294	R49555	R59556	AA009840	AA778675	AA629838	K39191	N6305/	***************************************	Walloo Laasaa	N32542	H22563	R54036	AA883187	AA047441	AA701315	T49355	N47012	AA676998	W81432	NOFZED	VA(92263	H79466	10/03717	T69270	AA418545	AA702335	H97440	R40434	44437142	R39951	AA609189	
6	960897	40880	151184	38344	41595	430092	1049033	884790	25,240	284714	300149	410034	270076	£1013 £1016	40010	1467161	488431	435099	67440	280155	454083	347586	269606	358000	235164	357373	82869	767315	447416	250069	28298	757389	25915	1031446	
;	87 (8 8	S 1	. 3	35	33	중 :	န္တ	8 l	34	3 6	n (} ;	÷	; ç	3 4	,	¥ 4	}	. 8	?	\$ \$	3 1	- £	4 5	3 2	; <u> </u>	3 4	3 2	5 4	3 8	8 8	3 8	. 29	

Table 2-11: Prostate Cancer Cell Lines vs. Normal Prostatic Epitheliaf Gell Lines. Genes upregulated 5 fold or more in at least 2 cancer cell lines.

ო	ო	ო	ო	ო	ო	ო	ო	ო	ო	ო	m	က	(n)	ო	က	m (m ·	m	ო	ო	ო	ო	ო	ო	က	ო	ო	හ -	ო	ო	ო	ო	ო	က	
5.35	5.12	20.74	15.62	17.41	22.12	19.45	8.12	6.86	11.26	7.37	24.06	17.09	16.72	20.94	12.92	7.26	5.03	15.10	7.37	5.79	11.50	7.50	9.62	7.35	5.24	5.84	13.62	5.71	6.90	10.11	11.70	5.53	9.84	6.83	
5.86	29.82	9.76	11.86	15.99	8.92	12.04	9.42	17.85	21.72	7.81	7.58	13.80	8.33	7.88	8.10	13.18	5.77	8.56	10.28	10.79	11.42	14.40	13.22	11.24	12.44	8.66	8.45	8.47	12.47	7.21	9.17	13.78	10.99	8.33	
31.21	6.39	10.64	13.47	7.33	9.24	8.23	22.16	14.72	5.65	21.95	5.26	5.65	10.15	5.06	11.07	11.48	20.21	5.69	11.04	11.98	5.53	6.47	5.48	9.36	10.06	12.99	5.38	13.21	7.38	9.14	5.30	69.9	5.02	9.78	
14.14	13.78	13.71	13.65	13.58	13.43	13.24	13.23	13.14	12.88	12.37	12.30	12.18	11.73	11.30	10.70	10.64	10.34	9.78	9.57	9.52	9.48	9.46	9.44	9.32	9.24	9.16	9.15	9.13	8.92	8.82	8.72	8.67	8.62	8.31	က
1.25	0.39	15.63	16.52	0.17	0.13	8.63	1.91	0.23	0.17	3.56	0.12	14.39	0.31	0.45	0.09	13.51	7.64	0.32	0.16	10.61	1.40	2.04	10.53	3,43	1.03	0.60	1.44	1.40	0.11	0.05	4.90	12.90	3.90	1.57	
60.0	0.03	1.14	121	0.01	0.01	0.65	0.14	0.02	0.0	0.29	0.01	1.18	0.03	0.0	0.01	1.27	0.74	0.03	0.02	1.12	0.15	0.22	1.12	0.37	0.11	0.07	0.16	0.15	0.01	0.01	0.56	1.49	0,45	0.19	
R42780	R42600	A 4628.867	AA508974	A4128617	A4016225	W/93379	AA885478	H19217	N34042	AA598531	AA644550	AA262211	AA609002	H67712	AA453433	H58234	AA678295	AA041197	AA701652	N47113	62206N	AA702684	N24042	AA045665	H04789	H77727	H46176	AA598817	W49672	T50699	AA187351	AA137072	AA520999	Re0019	
32109	30850	1035796	808286	502003	359269	415089	1466409	51103	244050	897807	845441	686172	1030798	210820	788209	204483	430928	376452	433567	280375	303089	448059	269300	489213	43733	234617	177884	897956	324901	78217	624627	491184	826273	42906	
ç	3 2	3 5	3 %	3 6	5 8	8 8	8 2		. 22	. E	74	75	76	1 2	78	6/	80	. <u> </u>	83	83	. 2	c	8 8	8 6	88	8	6	6	85	83	8	92	96	26	

Table 2-11: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes Lipregulated 5 fold or more in at least 2 cancer cell lines.

ო ო	ო	က	က	က	က	က	, ()	က	က	ო	က	ო	က	က	က	က	8	7	8	7	7	8	7	8	7	N	7	7	7	8	7	2	8		
8.51	9.05	76.7	11.50	7.97	9.46	8.78	7.98	5.19	5.60	6.63	5.79	5.08	5.21	5.52	5.77	5.40	7.96	7.05	1917.92	5.67	6.10	9.57	9.33	7.73	1.35	5.84	146.14	3.45	105.59	12.54	13.76	7.41	99.88		
11.10	6.02	77.5	7.09 7.10	6.65	6.4	7.63	7.83	5.36	5.02	5.78	5.67	5.73	6.93	7.41	5.63	503	2966.06	2506.90	64 14	1831.96	1516.35	1171.33	944 19	841.07	674.66	302.09	140.40	249.63	150.46	239.88	229.57	203.18	95.72		
5.27	11.86	0 ! 0 ! 1 0 !	7.7	4 6.		 	. v.	00 01	9.45	6.67	6.97	7 47	 	9 6		i, r	 	 0	08.1	2.32	 9	7 7		7:24	00.1	5.55 6.55		O. 60	- 00 00 00 00 00 00 00 00 00 00 00 00 00	50 C	D	2 40	5 C		
8.30	7.87	7.61	7.61	7.55	۲۶./	7.50	/.10 1	CD.,	0.00 0.00 0.00	000	0.0	5 6	0 6	6.03 1.00	9.69 9.00	5.63	5.20	991.80	838.64	660.46	613.07	508.28	393.88	319.36	283.26	228.86	103.67	95.78	86.03	85.65	85.17	81.58	71.20	8 4 .00 .	4
7.83	0.54	5.69	5.53	0.15	3.90	0.12	18.03	3.05	0.72	2.07	47.65	7.33	16.66	29.59	0.07	1.94	0.12	9.95	8.39	9.14	00.6 6	2.08	3.94	3.19	2.83	1.79	. 9.	1.93	2.66	6.83	0.51	0.82	0.71	5.16 9	
0.94	0.07	0.75	0.73	0.02	0.53	0.02	2.51	0.43	0.10	0.31	7.49	1.20	2.73	4.89	0.01	0.34	0.02	0.01	0.01	0.01	0.01	0.01	0.01	0.01	0.01	0.0	0.01	0.02	0.03	0.08	0.01	0.01	0.01	0.08	
AA425755	AA455938	R96941	AA608568	N30372	H29227	AA477400	AA663310	R06944	AA701645	R28287	H02231	R10284	AA425302	AA459318	R93875	H08785	R52085	N74995	AA009615	R60995	N57483	AA433885	N64379	AA099431	AA429904	AA446446	AA424900	H56453	AA071503	AA485214	AA459917	W80611	AA876021	AA419486	
773383	813271	200402	950690	260035	52704	740620	853368	126650	433576	134719	151055	128947	773073	814306	275871	45882	40299	299559	365551	42627	277327	773640	290213	510631	781157	781036	768226	203878	366039	815794	795627	347182	1161830	746152	
œ	9 9	5 5	5	102	103	\$	105	106	107	108	109	110	Ξ	112	113	414	115	116	417	. 4	119	120	121	122	123	124	125	126	127	128	129	130	131	132	

Table 2-11: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line Genes upregulated 5 fold or more in at least 2 cancer cell lines.

7	8	7	7	8	7	7	8	. 2	1 6	, ,	, ,	. 6	2	. ~	. 2	. 8	. ~		, ,	4 6	• •	, ,	4 6	4 (, ,		۰ ،	۱ ۸		4 (v (4 (7 (7	7	
11.76	0.73	4.30	70.45	123.70	0.72	1.93	Б	6 A7	42.47	<u> </u>	91.51	 	3 46	5 5	10.33	19:00	. c	. r. 5	27:16	6.63	- G	7.00	55.75	91.20	9 6	5.5	3, 1	1:15		52.45	1.00	63.66	10.31	9.84	6.05	
168.10	173.62	170.43	4.69	51 92	88 60	154 30	150.40	150.40	1.37	139.30	56.58	128.30	120.65	55.121	109.55	99.20	106.34	105.49	52.03	89.38	0.12	81.20	32.52 25. 1	7.97	76.38	48.0	70.25	69.04 40.03	1.36	27.83	72.00	11.75	66.04	62.71	68.60	
3.73	88	9 6	67.0	104:24	5.5	48.87 48.84	5.30	0.54	144.61	0.63	0.91	2.24	3.54	5.62	4.59	4.81	3.08	5.72	0.93	1.79	87.92	2.89	0.77	1.00	1.51	72.37	0.74	1.21	8.33	0.69	5.77	2.28	0.59	4.23	2.05	
61.20	17.04	60.41	50.34	59.79	58.88	56.02	53.90	52.11	50.82	50.72	49.67	45.99	43.48	43.37	41.72	41.11	38.18	37.79	34.73	32.46	31.22	30.56	30.35	30.08	27.65	27.63	27.34	27.13	27.04	26.99	26.26	25.90	25.65	25.59	25.57	ς.
5	5 6	0.83	1.60	26.93	2.87	12.41	0.28	2.21	60.94	1.10	13.02	09:0	0.81	0.43	0.42	0.41	0.38	0.38	8.56	1.12	2.62	0.70	7.89	0.30	5.88	16.16	0.27	0.27	0.20	14.38	0.26	14.15	0.39	0.29	1.90	
3	5.0	0.0	0.03	0.45	0.05	0.22	0.01	9.04	1.20	0.02	0.26	0.01	0.02	0.01	0.01	0.01	0.01	0.01	0.25	0.03	0.08	0.02	0.26	0.0	0.21	0.58	0.01	0.01	0.01	0.53	0.01	0.55	000	5 6	200	
	W94289	AA757754	AA461522	R23270	AA251784	W73144	AA086475	R40231	AA437224	R44741	AA504844	AA449120	AA855158	R37395	AA437126	AA810225	AA062985	AA857748	R91566	R59473	H29290	AA401434	AA485052	A A 0 9 3 6 9	R49439	R90934	R15740	AA460274	AA443624	H50655	AA425158	NAZAAS	CHANGE	192707	P402138	
	346942	395708	795858	131452	684879	344589	562811	30207	757435	33827	825740	785849	1389018	28498	757368	1367900	366209	1435339	196544	37823	49858	743189	815861	489656	38347	167205	53039	796549	771294	194318	769000	009086	660007	2023/8	77606	200
	133	<u>놓</u>	135	136	137	138	139	140	141		143	. 4	145	146	147	148	149	150	15.	15.		3 5	7.	3 4	157	52	159	160	£	5 6	701	3 3	\$ 5	2	5 5 7	<u> </u>

Table 2-11: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line, Genes upregulated 5 fold or more in at least 2 cancer cell lines.

7	7	7	7	7	7	7	7	7	. 2	7	8	7	~	. 2	۰ ۵	۰ ،		4 (٠ ,	v (v (7 (74 (71 (71 (74 (N (7 (N (7 (7	7	~	7	7		
16.63	5.46	8.05	11.62	31.35	45.50	11.25	62.76	43.13	3.46	48.25	25.34	31.76	11 14	K3 K1	64.20	23.52	2.00	50.5 50.5	47.62	17.48	28.27	20.17	50.17	45.41	25.26	49.74	46.16	1.08	43.06	5.54	10.71	6.22	5.54	16.22	8.77		
54.64	67.88	64.40	60.05	41.17	26.82	50.48	0 54	9.9. F. G.	60.63	23.5	13.13 13.64	42.07	5.00 m	51.85 Fž. 0	8.50 9.01	c0.01	1.50	23.72	10.88	42.76	30.54	38.72	7.58	10.31	30.81	4.0	1.93	14.31	10.12	47.06	38.83	45.91	45.44	34.15	40.61		
4.46	2.29	2.87	3.61	232	i (17.7	2. 6	4.0	0.95	8.08 10.08	0.87	00.L 90.	φ.4α Θ	3.82	2.34	0.88	39.83	37.52	4.40	0.93	1.29	1.00	1.10	2.79	2.35	7.38	9.69	41.41	3.39	2.02	4.49	1.21	1.87	1.97	2.37	i	
25.24	25.24	26.40	25.10	23.09	24.93	24.86	24.50	24.25	24.24	23.75	23.68	23.00	22.37	22.30	21.79	21.71	21.70	21.44	20.97	20.39	20.04	19.97	19.61	19.50	19.47	19.37	19.26	18.93	18.85	18.21	18.01	17.78	17.62	17.44	17.25	77.71	တ
200	0.23	0.23	0.25	0.17	0.25	24.91	0.24	26.54	7.41	0.84	3.51	0.23	0.16	3.74	0.34	3.65	0.14	40.77	19.16	6.19	3.94	0.20	3.26	0.20	3.30	0.19	0.19	21.01	0.17	0.74	0.35	3.45	, c		÷ 6	4	
Š		0.0	0.01	0.0	0.01	1.00	0.0	1.09	0.31	9.0	0.15	0.01	0.01	0.17	0.02	0.17	0.01	1.90	0.91	0:30	0.20	0.01	0.17	0.01	0.17	0.01	0.01	1.1	0.01	9.0	0.02	5 5	9 6	3 6	0.0	0.03	
1	AA010383	AA070437	R15813	AA236986	W94486	H19826	H96647	T57803	AA490945	R42112	AA504625	170198	AA630328	R44936	AA161097	AA491261	H10072	T61938	N32904	AA255954	R95684	A461508	AA520979	D86070	H28091	A 4 4 9 6 0 2 2	AA456628	T65736	R54492	AA620747	A A 0.57195	A 407FOF	20000	Kenozo	AA001376	AA489662	
!	430255	530958	53315	687972	357884	172517	251806	80643	824025	29585	825606	81050	855910	34204	591055	824270	46843	00062	259884	686552	199337	705834	826256	407265	162310	759163	809583	80338	39722	1049284	480078	409070	023040	42907	361899	823727	
	168	169	170	171	172	173	174	175	176	171	178	179	180	181	182	183	<u>\$</u>	185	186	187	, d	8 8	3 5	0 7	- 6	197	3 3	104	8 5	40.7	6 6	0 0	20 1	200	201	202	

Table 2-11: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 5 fold or more in at least 2 cancer cell lines.

64 (7	7	7	7	7	7	7	7	8	. 2		2		۰ ،	۰ ۵	,	4 C	۱ ,	7 (7	7	8	7	7	7	8	7	7	7	7	7	8	2	. 6	. 6	ı	
5.21	12.21	34.18	7.22	5.65	6.69	28.36	25.15	19.22	15.13	1 - 1 - 1 - 1 - 1 - 1 - 1 - 1 - 1 - 1 -	F. 87	15.54	20.00	23.23	28.45	6.15	53.10	09.72	84.4	12.18	18.27	29.19	18.18	7.18	26.50	7.37	25.70	28.39	2.23	3.21	5.10	0.58	5 6	 	20.03	7 0.90	
42.11	37.48	1.51	39.78	40.90	36 93	16.52	2.55	22.78	25.10	29.45	26.10	0.27	3.02	18.93	gr.4	9.24	36.28	1.04	18.97	26.44	21.58	8.82	18.08	29.16	45.	0.76	12.28	8.17	29.01	28.51	21 40	5 T T T		29:00	89°	9r.4r	
4.26	1.87	14.65		2. c	1 6		2.63	80.0°	4.69	1.53	0.42	36.64	24.69	1.00	10.07	5.00	0.88	2.99	17.85	2.42	0.64	2.08	3.39	808	10.97	30.80	30.00	, e	7 4	t a	6.51	55.7	22.57	7. .	10.40	1.96	
17.19	17 19	7 2	10.70	16.62	16.47	15.85	15.84	15.64	15.57	15.03	14.95	14.60	14.41	14.40	14.24	14.13	14.09	13.89	13.76	13.68	13.50	13.36	13.22		13. 4 4 CO	13.00	12.97	12.88	12.01	8/7ZL	12.67	12.64	12.56	12.41	12.39	12.38	7
98.0	90.6	97.	0.11	0.29	1.14	0.40	2.94	5.98	0:30	3.91	3.94	0.54	1.64	0.14	3.51	17.58	2.19	4.17	0.13	131	3 05		- 9c	0.20	 	0.08	0.36	0.13	8.37	0.71	0.25	1.99	0.13	4.99	9.44	0.12	
9	9 6	90.0	0.01	0.02	0.07	0.03	0.19	0.38	0.02	0.26	0.26	9.0	0.11	0.01	0.25	1.24	0.16	0:30	5) - C	6.60	0.03	0.02	0.08	0.01	0.03	0.01	0.65	90.0	0.05	0.16	0.01	0.40	0.76	0.01	
	AA609304	AA857163	AA481481	N26559	N51297	AA410437	AA504838	AA417713	AA045175	AA463445	H18932	AA035745	AA476543	AA434388	AA883800	AA774665	A A 699601	W90705	2000	K56149	K3//30	AA427719	H15549	AA491227	AA883181	N52193	AA035730	AA017379	AA911236	R43915	AA621355	AA070435	AA453616	AA437212	AA620697	AA022496	
	1031568	1410444	756629	266366	283070	753411	825726	746232	487766	811766	51599	364865	785293	770860	1461609	856789	436402	418004	10001	41406	26806	771004	49499	824074	1467166	284288	364844	361122	1526789	33066	1055121	530954	795446	757428	1049185	364352	
	203	5 8	202	506	207	208	508	210	,	212		214	. 4	2.5	24.5	: č	2 6	617	37	224	222	223	554	225	226	227	228	229	230	231	232	233	234	235	236	237	

0 0	, ,		۱ ،	. 6	8	. 2	. 2	۱	. ~	. 6	8	8	2	7	7	7	2	8	7	2	7	7	7	23	7	8	7	7	7	8	7	2	8		
11.09	3 1	17.55	23.17	4.0.10 80 80 80 80	16.24	10:11		0.00	5 ° 6	\$ 60 CC	24.11	16.34	7.83	13.80	4.94	16.62	10.20	12.02	20.10	8.36	5.22	21.77	5.14	6.78	15.60	23.76	15.64	0.48	1.65	17.42	21.42	5.78	1.12		
23.87	29.43	15.47	4.23	14.95	18.70	51.71	13.18	20.54	19.91	20.61	10.16	5. 5. 7. 8.	22.78	17.96	58	15.37	20.21	15.78	10:10	00	22.31	8.99	4.33	21.63	14.48	6.31	14.23	7.21	17.15	11.85	909	20.48	20.18		
2.11	6.29	3.28	8.77	1.00	4. 4.	2.08	3.09	0.58	3.60	10.87	3.65	86.4	55.0	9 6	0.92	7.7.	- 60 6	2.03		50.7 50 66	14.3	44.0	2.5	267		0 90 C	22.0	22.87	14 54	10.11	0.93	2.01	, a	- - -	
12.36	12.24	12.10	12.06	12.02	11.98	11.82	11.80	11.71	11.67	11.63	11.61	11.46	11.17	11.06	10.90	10.87	10.84	10.83	10.81	10.79	10.76	10.66	10.30	10.45	10.36	10.32	10.31	10.21	er.01	10.11	10.07	10.05	70.00 10.00)B.D	∞
3.72	0.12	0.31	16.58	1.69	16.82	23.53	3.89	0.18	11.24	0.20	27.64	0.11	0.34	0.95	3.90	5.68	6.12	0.31	90.0	9.32	0.11	5.93	8.09	12.13	0.44	0.87	4.21	5.55	9.47	0.12	2.53	16.05	0.37	2.42	
0:30	0.01	0.03	88.	0.14	1.40	1.99	0.33	0.02	0.96	0.02	2.38	0.01	0.03	0.09	0.36	0.52	0.57	0.03	0.01	0.86	0.01	0.56	0.77	1.16	9. 8	0.08	0.41	0.54	0.93	0.01	0.25	1.60	9.0	0.24	
H10713	167927	15/32/	AA476576	R92201	AA706901	AA477227	H09940	AA161188	R44816	W93523	AA026682	AA179161	AA160670	AA427737	H48070	AA460299	AA410434	N74131	AA461513	N30751	AA680136	AA417920	AA489073	AA454949	N93721	AA281784	H54451	N49389	AA412049	AA158035	R94504	AA478036	AA598787	AA626167	
47764	11674	1,017	101010	405845	451649	739450	46461	591157	33839	357138	366971	609228	592728	770854	193333	795746	753428	298417	795840	257955	433155	767403	824802	814769	307138	712401	203184	277621	729953	590640	197651	740457	898073	745248	
o C	8	88 C	047	L 47	743	? ?	¥ ;	245	2 17 6	748 248	249	250	251	252	253	254	255	256	257	258	259	260	261	262	263	264	265	266	267	268	269	270	271	272	

Table 2-11: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelia Cell Lines Genes upregulated 5 fold or more in at least 2 cancer cell lines.

0.50 4.94 0.39 3.82 0.001 0.09 0.01 0.09 0.02 0.21 0.02 0.21 0.02 0.21 0.02 0.21 0.02 0.01 0.03 0.04 0.01 0.09 0.01 0.09 0.01 0.09 0.01 0.09 0.01 0.09 0.01 0.09 0.01 0.09 0.01 0.09 0.01 0.09 0.01 0.09 0.01 0.09 0.01 0.09 0.02 0.01 0.03 0.02 0.03 0.26 0.03 0.26 0.03 0.26	855391 A	AA664007 AA625979	0.23	2.29	9.95 9.95	1.40	3.07	16.30 14.06 15.31	0 0 0
0.39 0.38 0.382 0.387 0.01 0.009 0.986 0.02 0.21 0.02 0.21 0.02 0.21 0.03 0.04 0.05 0.05 0.05 0.05 0.05 0.05 0.07 0.01 0.00 0.03 0.04 0.00 0.00 0.00 0.00 0.00	- ≶	292226	0.50	4.94	9.89	0.67	13.69	10.88	1 8
0.00 0.00 5.59 23.34 0.00 0.57 9.79 3.40 9.94 0.00 0.57 9.79 3.40 9.94 0.02 0.21 9.77 1.35 22.17 0.02 0.17 9.68 1.31 9.16 0.02 0.17 9.64 4.99 9.81 0.02 0.17 9.64 4.99 9.81 0.02 0.17 9.62 0.66 21.91 0.04 6.12 9.62 0.66 21.91 0.05 0.16 9.62 0.66 21.91 0.07 9.62 3.83 6.58 0.01 0.10 9.62 3.83 6.58 0.02 0.16 9.62 3.83 6.58 0.17 9.64 4.99 9.81 1.61 0.17 9.53 1.61 1.75 0.16 9.47 10.45 1.04 0.17 0.09 <	-	V21633	0.39	3.82	9.87 9.86	8.07	19.71	1.81	7
0.06 0.57 9.79 3.40 9.94 0.02 0.21 9.77 1.35 22.17 1.26 0.21 9.77 1.35 22.17 0.62 5.97 9.68 1.31 9.16 0.02 0.17 9.64 4.99 9.81 0.02 0.17 9.62 0.66 21.91 0.03 0.10 9.62 0.66 21.91 0.04 0.10 9.62 3.83 6.58 0.01 0.10 9.62 3.83 6.58 0.02 0.16 9.62 0.66 21.91 0.03 0.77 9.48 1.15 11.85 0.04 0.09 9.40 8.70 1.04 0.07 0.49 8.74 1.04 5.97 0.07 0.09 9.40 8.70 1.04 0.01 0.09 9.40 8.70 1.04 0.02 5.85 9.41 4.56	•	R55220	20.0	0.03	9.80	5.59	23.34	0.46	0 0
0.02 0.21 9.77 1.35 22.17 1.26 9.74 3.45 18.90 0.62 5.97 9.68 1.31 9.16 0.02 0.17 9.64 4.99 9.81 0.02 0.17 9.62 0.66 21.91 0.04 6.12 9.62 0.66 21.91 0.05 0.10 9.62 0.66 21.91 0.07 0.10 9.62 0.66 21.91 0.08 0.77 9.53 2.05 1.50 0.15 1.40 9.48 1.15 11.85 0.15 1.40 9.48 1.15 11.85 0.15 1.40 9.48 1.15 11.85 0.07 7.14 9.41 4.56 7.97 0.07 5.02 9.39 1.84 5.32 0.01 0.08 9.40 8.70 1.04 0.02 0.19 9.36 8.59 1.12		41003636	90.0	0.57	9.79	3.40	9.94	16.03	4 (
1.26 9.74 3.45 18.90 0.62 5.97 9.68 1.31 9.16 0.02 0.17 9.64 4.99 9.81 0.02 0.17 9.62 0.66 21.91 0.04 6.12 9.62 0.66 21.91 0.07 9.62 3.83 6.58 0.01 0.10 9.62 3.83 6.58 0.02 0.16 9.62 3.83 6.58 0.03 0.77 9.53 2.05 1.563 0.04 0.77 9.48 1.15 11.85 0.19 9.47 1.045 1.00 0.76 7.14 9.41 4.56 7.97 0.62 5.85 9.41 4.56 7.97 0.07 0.09 9.40 8.70 1.04 0.01 0.08 9.40 8.70 1.14 0.01 0.08 9.39 1.84 5.37 0.01 <		A120779	0.02	0.21	9.77	1.35	22.17	9. (8 87	۸ ۷
0.62 5.97 9.68 1.31 9.10 0.02 0.17 9.64 4.99 9.81 0.04 6.12 9.62 0.66 21.91 0.04 6.12 9.62 0.66 21.91 0.02 0.16 9.62 3.83 6.58 0.02 0.16 9.56 1.61 17.50 0.08 0.77 9.53 2.05 15.63 0.09 0.77 9.48 1.15 11.85 0.15 1.40 9.48 1.15 11.85 0.15 1.40 9.47 10.45 1.00 0.76 7.14 9.41 4.56 7.97 0.62 5.85 9.41 4.56 7.97 0.07 0.09 9.40 8.70 1.04 0.01 0.09 9.40 8.70 1.04 0.02 5.02 9.39 1.84 5.37 0.03 0.22 9.31 1.34		A682533	1.26	12.26	9.74	3.45	18.90	18.56	1 74
0.02 0.17 9.64 4.39 2.131 0.64 6.12 9.62 0.66 21.91 0.01 0.10 9.62 0.66 21.91 0.02 0.16 9.56 1.61 17.50 0.08 0.77 9.53 2.05 15.63 0.09 0.77 9.48 1.15 11.85 0.01 0.09 9.47 10.45 1.00 0.76 7.14 9.41 4.56 7.97 0.05 5.85 9.40 8.70 1.04 0.07 0.09 9.40 8.70 1.04 0.01 0.09 9.40 8.70 1.04 0.01 0.09 9.40 8.70 1.04 0.01 0.09 9.40 8.70 1.04 0.02 9.39 1.84 5.32 0.01 0.08 9.36 8.59 1.12 0.02 0.19 9.32 1.34 16.72 0.10 0.09 9.20 1.34 10.86 0.1		4A620528	0.62	5.97	9.68	1.31	9. 9. 5. 8. 9.	14.13	8
0.64 6.12 9.62 0.00 6.18 0.01 0.10 9.62 3.83 6.58 0.02 0.16 9.56 1.61 17.50 0.08 0.77 9.53 2.05 15.63 0.09 0.47 1.15 11.85 1.00 0.01 0.09 9.47 10.45 1.00 0.76 7.14 9.41 4.56 7.97 0.62 5.85 9.41 4.56 7.97 0.01 0.09 9.40 8.70 1.04 0.01 0.09 9.40 8.70 1.04 0.01 0.09 9.40 8.70 1.04 0.01 0.08 9.36 8.59 1.12 0.01 0.06 9.34 11.43 1.54 0.02 0.19 9.32 3.79 1.55 0.02 0.19 9.22 2.30 7.70 0.44 4.10 9.26 2.30 7.70 0.44 4.10 9.26 2.30 7.70 <t< td=""><td></td><td>AA157261</td><td>0.02</td><td>0.17</td><td>9.0 49.0</td><td>יים מיים מיים</td><td>2. 9.</td><td>6.28</td><td>8</td></t<>		AA157261	0.02	0.17	9.0 49.0	יים מיים מיים	2. 9.	6.28	8
0.01 0.10 9.62 0.00 0.00 0.02 0.16 9.56 1.61 17.50 0.08 0.77 9.53 2.05 15.63 0.15 1.40 9.48 1.15 11.85 0.01 0.09 9.47 10.45 1.00 0.01 0.09 9.47 10.45 1.00 0.01 0.09 9.40 8.70 1.04 0.01 0.09 9.36 8.59 1.12 0.01 0.06 9.36 8.59 1.12 0.02 0.22 9.39 1.84 5.32 0.02 0.22 9.39 1.84 5.32 0.02 0.22 9.31 0.97 12.53 0.03 0.26 8.35 9.29 1.35 16.72 0.44 4.10 9.26 2.30 7.70 0.44 4.10 9.26 2.30 7.70 0.40 0.97 9.22 12.49 10.86 0.10 0.87 9.18 2.37 19.73 0.03 0.26 9.09 3.46 10.73 0.03 0.26 9.09 3.46 10.73 0.03 0.26 9.07 12.50 0.01 0.09 9.07 12.50 0.36 0.01 0.09 9.07 5.08 1.00		AA133590	0.64	6.12	9.62	0 c	5.58	18.44	8
0.02 0.16 9.56 1.61 17.00 0.08 0.77 9.53 2.05 15.63 0.15 1.40 9.48 1.15 11.85 0.01 0.09 9.47 10.45 1.00 0.01 0.09 9.47 10.45 1.00 0.05 5.85 9.41 4.56 7.97 0.01 0.09 9.40 8.70 1.04 0.01 0.09 9.40 8.70 1.04 0.01 0.09 9.40 8.70 1.04 0.01 0.09 9.30 1.84 5.32 0.01 0.08 9.36 8.59 1.12 0.01 0.06 9.34 11.43 1.54 0.02 0.19 9.32 3.79 15.51 0.02 0.19 9.22 1.35 16.72 0.44 4.10 9.26 2.30 7.70 0.44 4.10 9.26 2.30 7.70 0.10 0.87 9.18 2.37 19.73	•	AA156342	0.01	0.10	9.62	20.0	47.50	60 60	8
0.08 0.77 9.53 2.05 15.05 0.15 0.15 0.15 0.15 0.15 0.15		R89490	0.02	0.16	9.56	1.61	17.50	10.89	. 7
0.15		R46000	0.08	0.77	9.53	2.05	15.63	15.03	8
0.01 0.09 9.47 10.45 1.00 0.76 7.14 9.41 3.48 5.97 0.62 5.85 9.41 4.56 7.97 0.62 0.03 9.40 8.70 1.04 0.01 0.09 9.40 8.70 1.04 0.01 0.08 9.36 8.59 1.12 0.01 0.06 9.34 11.43 1.54 0.02 0.19 9.32 3.79 15.51 0.02 0.44 4.10 9.26 2.30 7.70 0.44 4.10 9.26 2.30 7.70 0.40 0.40 0.97 9.22 12.49 10.86 0.10 0.87 9.12 0.81 11.63 0.35 3.21 9.15 8.17 14.83 0.35 0.03 0.26 9.09 3.46 10.73 0.03 0.26 9.07 12.50 0.36 0.00 0.00 9.07 5.08 1.00 0.00 0.00 9.07 5.08 1.00 0.00 0.00 9.07 5.08 1.00	-	AA219060	0.15	1.40	9.48	1.15		16.97	. 7
0.76 7.14 9.41 3.48 9.97 0.62 5.85 9.41 4.56 7.97 0.62 5.85 9.41 4.56 7.97 0.01 0.09 9.40 8.70 1.04 0.01 0.08 9.36 8.59 1.12 0.01 0.06 9.34 11.43 1.54 0.02 0.19 9.32 3.79 1.54 0.02 0.19 9.32 3.79 15.51 0.02 0.22 9.31 0.97 12.53 0.08 6.35 9.29 1.35 16.72 0.44 4.10 9.26 2.30 7.70 0.44 4.10 9.26 2.30 7.70 0.10 0.87 9.18 2.37 19.73 0.10 0.87 9.15 8.17 14.83 0.24 9.15 9.15 0.81 10.73 0.03 0.26 9.09 3.46 1.00 0.01 0.09 9.07 5.08 1.00 </td <td></td> <td>W16659</td> <td>0.01</td> <td>60:0</td> <td>9.47</td> <td>10.45</td> <td>00.r</td> <td>18 70</td> <td>. 0</td>		W16659	0.01	60:0	9.47	10.45	00.r	18 70	. 0
0.62 5.85 9.41 4.56 7.97 0.01 0.09 9.40 8.70 1.04 0.053 5.02 9.39 1.84 5.32 0.01 0.08 9.36 8.59 1.12 0.01 0.06 9.34 11.43 1.54 0.02 0.19 9.32 3.79 1.54 0.02 0.22 9.31 0.97 12.53 0.02 0.22 9.29 1.35 16.72 0.44 4.10 9.26 2.30 7.70 0.44 4.10 9.26 2.30 7.70 0.10 0.87 9.25 1.35 10.86 0.10 0.87 9.18 2.37 19.73 1.08 9.89 9.17 4.58 11.63 0.27 2.44 9.15 8.17 14.83 0.27 2.44 9.15 0.81 10.73 0.03 0.26 9.09 3.46 10.73 0.01 0.02 9.07 12.50 0.36	•	AA678348	0.76	7.14	9.41	3.48	/6.c	10.73	. ~
0.01 0.09 9.40 8.70 1.04 0.53 5.02 9.39 1.84 5.32 0.01 0.08 9.36 8.59 1.12 0.01 0.06 9.34 11.43 1.54 0.02 0.19 9.32 3.79 1.54 0.02 0.22 9.31 0.97 12.53 0.68 6.35 9.29 1.35 16.72 0.44 4.10 9.26 2.30 7.70 0.44 4.10 9.26 2.30 7.70 0.10 0.87 9.22 12.49 10.86 0.10 0.87 9.18 2.37 19.73 1.08 9.89 9.17 4.58 11.63 0.27 2.44 9.15 8.17 14.83 0.27 2.44 9.15 8.17 14.83 0.03 0.26 9.09 3.46 10.73 0.03 0.26 9.07 5.08 1.00 0.01 0.09 9.07 5.08 1.00 0.01 0.09 9.07 5.08 1.00		AA204830	0.62	5.85	9.41	4.56	76.7		۰ ۱
0.53 5.02 9.39 1.84 5.32 0.01 0.08 9.36 8.59 1.12 0.01 0.06 9.34 11.43 1.54 0.02 0.19 9.32 3.79 15.51 0.02 0.22 9.31 0.97 15.51 0.02 0.22 9.29 1.35 16.72 0.68 6.35 9.29 1.35 16.72 0.44 4.10 9.26 2.30 7.70 0.44 4.10 9.26 2.30 7.70 0.10 0.87 9.22 12.49 10.86 0.10 0.87 9.18 2.37 19.73 1.08 9.89 9.17 4.58 11.63 0.27 2.44 9.15 8.17 14.83 0.03 0.26 9.09 3.46 10.73 0.03 0.26 9.07 5.08 1.00 0.01 0.09 9.07 5.08 1.00		AA679180	0.01	0.09	9.40	8.70	40.	5.50	۰ ۱
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0.01 0.06 9.34 11.43 1.54 0.02 0.19 9.32 3.79 15.51 0.02 0.22 9.31 0.97 12.53 0.08 6.35 9.29 1.35 16.72 0.44 4.10 9.26 2.30 7.70 0.44 4.10 9.26 2.30 7.70 0.10 0.87 9.18 2.37 19.73 1.08 9.89 9.17 4.58 11.63 0.35 3.21 9.15 8.17 14.83 0.27 2.44 9.15 8.17 14.83 0.03 0.26 9.09 3.46 10.73 0.01 0.09 9.07 12.50 0.36 0.01 0.09 9.07 5.08 1.00		AA699926	0.01	0.08	9.36	8.59	21.1		
0.02 0.19 9.32 3.79 15.51 0.02 0.22 9.31 0.97 12.53 0.06 6.35 9.29 1.35 16.72 0.44 4.10 9.26 2.30 7.70 0.10 0.97 9.26 2.30 7.70 0.10 0.87 9.18 12.49 10.86 0.10 0.87 9.18 2.37 19.73 1.08 9.89 9.17 4.58 11.63 0.35 3.21 9.15 8.17 14.83 0.27 2.44 9.12 0.81 10.73 0.03 0.26 9.09 3.46 10.73 0.01 0.09 9.07 5.08 1.00		AA284072	0.01	90.0	9.34	11.43	<u>.</u> ;	0.00	۱ ۵
0.02 0.31 0.97 12.53 0.68 6.35 9.29 1.35 16.72 0.44 4.10 9.26 2.30 7.70 0.44 4.10 9.26 2.30 7.70 0.10 0.87 9.18 2.37 19.73 0.10 0.87 9.18 2.37 19.73 1.08 9.89 9.17 4.58 11.63 0.35 3.21 9.15 8.17 14.83 0.27 2.44 9.12 0.81 13.04 0.03 0.26 9.09 3.46 10.73 0.03 0.26 9.07 12.50 0.36 0.01 0.09 9.07 5.08 1.00		W74377	0.02	0.19	9.32	3.79	15.51	9.0	, ,
0.68 6.35 9.29 1.35 16.72 0.44 4.10 9.26 2.30 7.70 0.10 0.97 9.22 12.49 10.86 0.10 0.87 9.18 2.37 19.73 1.08 9.89 9.17 4.58 11.63 0.35 3.21 9.15 8.17 14.83 0.27 2.44 9.12 0.81 13.04 0.03 0.26 9.09 3.46 10.73 0.03 0.26 9.07 12.50 0.36 0.01 0.09 9.07 5.08 1.00		N50962	0.02	0.22	9.31	0.97	12.53	44.44	, ,
0.44 4.10 9.26 2.30 7.70 0.10 0.97 9.22 12.49 10.86 0.10 0.87 9.18 2.37 19.73 1.08 9.89 9.17 4.58 11.63 0.35 3.21 9.15 8.17 14.83 0.27 2.44 9.12 0.81 13.04 0.03 0.26 9.09 3.46 10.73 0.03 0.26 9.07 12.50 0.36		AA455242	0.68	6.35	9.29	1.35	16.72	9.79	۰ ،
0.10 0.97 9.22 12.49 10.86 0.10 0.87 9.18 2.37 19.73 1.08 9.89 9.17 4.58 11.63 0.35 3.21 9.15 8.17 14.83 0.27 2.44 9.12 0.81 13.04 0.03 0.26 9.09 3.46 10.73 0.03 0.26 9.07 12.50 0.36		A1017607	44.0	4.10	9.26	2.30	0.70		, ,
0.10 0.87 9.18 2.37 19.73 1.08 9.89 9.17 4.58 11.63 0.35 3.21 9.15 8.17 14.83 0.27 2.44 9.12 0.81 13.04 0.03 0.26 9.09 3.46 10.73 0.03 0.26 9.07 12.50 0.36		AA705047	0.10	0.97	9.22	12.49	10.86	کن.4 د د د	4 (
1.08 9.89 9.17 4.58 11.63 0.35 3.21 9.15 8.17 14.83 0.27 2.44 9.12 0.81 13.04 0.03 0.26 9.07 12.50 0.36 0.01 0.09 9.07 5.08 1.00		D05067	0.10	0.87	9.18	2.37	19.73	5.45	v
0.35 3.21 9.15 8.17 14.83 0.27 2.44 9.12 0.81 13.04 0.03 0.26 9.09 3.46 10.73 0.03 0.26 9.07 12.50 0.36 0.01 0.09 9.07 5.08 1.00		206067	9 -	68.6	9.17	4.58	11.63	11.30	74 (
0.27 2.44 9.12 0.81 13.04 0.03 0.26 9.09 3.46 10.73 0.03 0.26 9.07 12.50 0.36 0.01 0.09 9.07 5.08 1.00		A446462	38.0	3.21	9.15	8.17	14.83	4.46	71
0.03 0.26 9.09 3.46 10.73 0.03 0.26 9.07 12.50 0.36 0.01 0.09 9.07 5.08 1.00		701010	72.0	2.44	9.12	0.81	13.04	13.51	7
0.03 0.26 9.07 12.50 0.36 0.01 0.09 9.07 5.08 1.00		13057 B	; O	0.26	60.6	3.46	10.73	13.08	7
0.01 0.09 9.07 5.08 1.00		K69364	8 6	0.26	9.07	12.50	0.36	14.35	7
0000		W44/00	8 6	60 0	9.07	5.08	1.00	21.12	8
		AA1000so	- - -	<u>;</u>	σ				

Table 2-11: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelia Cell Line: Genes upregulated 5 fold or more in at least 2 cancer cell lines.

7	7	7	73	7	7	7	8	2	7	- 7	. 2	. 2		. ^	۰ ،	• •	4 (4 (4 (N (7 (7	7	64	7	7	84	7	7	7	7	8	، ۱	4 (N	7	
14.67	5.38	9.46	1.10	16.60	5.40	11.94	8.45	14.43	08.80	18 10	15.48	7.50	35.1.4 AC 1.4	4. 5. 6. 8. 6.	800	3.90	98.7.	11.2/	12.36	17.21	6.37	14.40	16.24	19.27	2.96	13.09	11.26	18.07	7.71	14.93	15.23	78	o e t c	o :	13.11	13.46	
10.87	20.24	15.22	13.45	0.30	20.84	2.96	47 4R	2 80	27.70	¥ 70 0	† 6	0.03	00:01	1.00	75.0	14.21	6.50	12.46	11.51	6.16	13.73	6.65	6.27	0.41	13.18	1.67	11.14	0.56	13.64	, t	· ·	08.0	13.04	18.34	7.53	8.34	
1.65	1.54	2.44	12.54	10.05	200	5 5	7	0.92	9.82	0.36	1.34	3.86	1.87	10.33	6.22	7.42	0.78	1.45	1.30	1.81	4.95	3.89	2.41	5.22	8.74	86	2.23	5.83	200	t 600	0. (c)	9. 9. 9.	1.22	0.38	3.42	2.25	
9.06	905	9 04		5 6	0 0	Ø. 60	8.87	8.84	8.81	8.70	8.69	8.66	8.65	8.56	8.53	8.51	8.42	8.39	8.39	8.39	8.35	8.32	8.31	8.30	8.29	. a	9.£3	α 1 τ 1 τ		8.13 5.13	8.13	8.09	8.08	8.03	8.02	8.02	10
3.21	3.67	10.62	70.07	17.11	17.81	0.21	5.97	3.32	2.16	39.21	0.80	48.92	34.28	0.09	5.11	0.14	6.98	9.51	3.36	0.08	0.19	4.39	13.45	10.75	2.5	- c		n •		0.25	21.34	10.44	6.11	0.22	0.28	0.08	
3,5	3 6	9 1	1.17	1.30	1.98	0.02	0.67	0.38	0.25	4.51	0.09	5.65	3.96	0.01	09.0	0.02	0.83	1.13	0.40	0.01	0.02	0.53	1 63	<u> </u>	 	5 6	0.02	5. U	0.02	0.03	2.63	1.29	0.76	0.03	0.03	0.0	
7	C1 C07N	AA192527	AA488175	AA172056	AA620485	AA416970	W68220	H41203	H16772	AA629999	AA778646	H95638	AA775863	AA496998	AA682321	T52311	N80764	W47364	AA504265	AA036974	AA486265	AA465205	M4507.00	N/401/	AA488324	R61372	N50806	AA058576	AA131239	W96216	R41329	H54752	AA699427	AA457576	A A 705 825	H04044	
	667997	627541	877632	594758	951241	730036	342640	175533	50227	884511	1048995	242952	878544	897546	462926	71977	300615	324717	825418	484535	40450	800557	/66800	296429	842968	37980	280882	380884	503581	358599	30148	203275	433253	838776	4202182	230281	
	308	309	310	311	312	313	314	315	316	317	318	319	320	321	322	33	327	32 5	226	327	770	328	875	330	331	332	333	334	335	336	337	338	339	340	} ;		ţ

Table 2-11: Prostate Cancer Cell Lines vs. Normal Prostatic Epithefial Cell Lines Genes upregulated 5 fold or more in at least 2 cancer cell lines.

ç	00000	0000		8 8 18 8	00 8	0.92	14.68	8.40	8
3 3	195097	02/102	2	000	68 2	1.06	13.94	8.68	~
\$;	67910/	AA264112	67.0	2.5 2.0	7.83	2.27	15.21	6.01	7
£ .	38833	K507/5	0 6	0 0	7.77	4.86	11.97	6.49	8
346	190325	H29897) () ()	6.00 6.00 6.00 6.00	77.7	3 94	9.24	10.12	8
74	745393	AA625764	0.08 1	0.00		7 5 7	7.74	4.46	8
348	782259	AA431741	0.75	5.81	07.7		7.43	15.20	7
349	502444	AA156795	0.02	41.0	7.73	4 	8.93	10.00	8
350	91900cr	AA906257	5.0	24 59	7.70	7.55	4.17	11.39	2
501	107171	190710	5.5	13.42	7.70	4.60	7.66	10.84	7
352	204644	15560/ H57082	0.27	2.05	7.70	1.18	12.63	9.28	8
35.	322461	107.505	0 22	1.68	7.70	6.23	3.58	13.28	7
3 5	810881	A 4 63082	6	0.08	7.69	7.96	14.11	1.00	7
25.	882510	AA876460	34 04	261.61	7.68	3.35	8.00	11.71	7
3, 75	46977	H10234	200	0.53	7.67	2.09	11.89	9.03	7
35.8	46740	H10068	0 03	0.25	7.66	1.41	14.72	6.84	7
3,50	1035664	44780365	0.13	1.03	7.65	12.36	8.59	2.00	7
360	345743	W72666	2.36	18.06	7.64	1.23	16.02	5.66	7
361	120695	TOSBUG	000	0.08	7.61	0.83	12.70	9.32	74
. cg	725340	44291773	0.72	5.46	7.60	1.27	5.74	15.79	73
383	530820	AA070331	0.01	90.0	7.55	10.53	4.96	7.15	7
38	448032	AA702802	0.01	0.08	7.54	5.64	1.00	15.99	7
365	39833	R53455	26.0	7.28	7.53	0.35	12.36	9.88	7
386	770879	AA434403	0.75	5.64	7.53	3.47	10.88	8.23	7
367	868169	AA633835	0.16	1.18	7.49	1.25	5.39	15.83	~
368	811166	AA485749	0.01	0.08	7.46	12.26	0.92	9.18	7
369	83666	T61122	0.03	0.20	7.44	12.64	4.41	5.28	7
370	128243	R12473	0.83	6.14	7.44	1.30	8.97	12.04	7
371	37538	R49644	9.0	0.31	7.41	0.99	11.96	9.29	64
372	435076	AA701455	2.95	21.88	7.41	10.35	6.94	4.93	~
373	309288	N93924	3.23	23.90	7.39	5.03	4.68	12.46	7
374	843163	AA488367	7.80	57.27	7.34	3.00	6.84	12.18	63
375	278729	N62936	1.35	9.91	7.34	0.34	13.66	8.02	61
376	66317	T66816	3.88	28.27	7.29	6.83	0.63	14.40	7
377	451907	AA706968	2.08	15.11	7.27	11.07	0.08	10.67	73
					7				

7	N	7	7	2	7	84	8	7	7	8	7	7	8	7	7	7	7	8	` 7	74	7	7	7	7	7	7	7	7	8	~	7	7	7	8		
10.22	12.41	6.82	6.11	10.54	6.43	5.38	66.	11.27	10.38	10.69	11.80	7.84	7.62	5.32	8.84	6.81	3.65	6.50	11.87	9.20	14.49	11.35	9.27	10.38	1,16	3.14	9.86	5.69	8.65	12.53	8.19	9.34	11.61	5.09		
9.92	2.06	13.06	12.65	7 03	13.46	1 9	S::-	9 7	10.7	, c	8.75	0.03	11.28	2.34	11.35	12.50	5.87	9 9	7.50	70.7	7.67		9 22	88	7 T	11.26	4.41	10.34	9.72	1.10	151	8 48		7 t	3	
1.57	7.23			76.7	- 6	0.70	9.39 9.00	3.01 	2.72	67.7	04.4	42.67	2.5.5	7:4 13 68	9 9 7	<u></u>	3 ;	7/11	0.4	8.5	4. C	9 6	9.00	5.5	2.15	14.50 0.40	649		, t	; ; ;	7.7	10.80 F C	5.7). []	7 C.1	
7.24	7.23	7. 1	27.7	7.23	7.23	7.22	7.22	7.20	7.20	7.17	7.15	7.14	2.13	21.7	11.7	7.11	7.11	7.08	7.03	7.02	7.00	7.00	6.96 6.96	6.85	6.85	6.93	0 9. 0 0	0.92	76.9 76.0	0 0	Ö.Ö	6.88	6.85	6.84	6.83	72
35.0	3 6	7.70	1.17	8.99	5.80	0.31	33.83	6.75	4.64	15.96	0.10	0.11	105.24	14.44	0.62	1.51	95.95	14.78	0.08	2.01	100.32	0.28	0.44	0.68	36.26	0.07	0.39	3.69	7.46	3.78	90.0	2.74	4.75	0.83	0.25	
ğ	6.03	0.30	0.16	1.24	0.80	9.0	4.68	0.94	0.64	2.23	0.01	0.02	14.77	2.03	0.09	0.21	13.50	5.09	0.01	0.29	14.33	9.0	90.0	0.10	5.22	0.01	90.0	0.53	1.08	0.55	0.01	0.40	0.69	0.12	9.0	
	N22766	R43308	H19234	AA135809	T62072	AA011182	H97488	AA460952	AA455521	AA504201	AA456611	AA677655	AA894557	AA455237	AA449118	R96520	N66158	N45236	R74203	N53940	AI015359	W72556	W86196	T68878	AA485453	T58146	R56046	N20305	AA465238	AA886739	AA669222	AA452278	AA129135	AI025259	AA398406	
	266631	32756	50904	565734	85670	359722	251936	796114	809828	825282	809567	897271	1416782	814779	785845	199644	278531	283173	143365	247050	1637343	345330	415906	82215	811067	79254	40773	264502	814225	1500162	856585	787938	586650	1641367	726791	
	378	379	380	381	382	383	384	38.	386	387	388	389	390	391	392	393	394	395	396	397	398	399	9	104	405	403	4	405	406	407	408	409	410	11	412	•

Table 2-11: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line Genes upregulated 5 fold or more in at least 2 cancer cell lines.

7	7	7	7	~	7	8	•		۰,		4 (4 (7	7	7	7	7	7	7	۰ ،	4 (4 (v	7	7	7	7	7	8	7	7	, ,	4 (4 (7	7	7		
12.23	6.48	5.38	6.30	4.64	10.97	88.0		0.00	6.	4 (F	7.33	6.95	8.44	7.62	8.09	8.12	1.85	9.54	4.80) ()	0 0	3.05	4 .63	5.67	6.74	5.55	6.04	10.01	7.59	76.7	8 78) i	φ. (c)	6.08	9.70	9.25	1.28		
5.03	12.71	10.24	4.78	934	3.78	0 1 0	67.7	1.82	4.68	8.01	10.73	9.66	4.39	4.47	0.58	3.71	10.05	9 18	2 3	44.7	7.59	5.69	9.69	8.97	11.33	11.59	2.65	5.16	9.29	8 32	0.32	2 5	5.78	11.14	6.70	4.97	12.25		
3.20	1.22	4.75	0.00	9:50	5 1	5.5/	12.08	9.56	7.84	8.02	2.08	3.37	7.13	7.85	11.27	8	4 60	10. T	01.T	7.52	2.44	10.89	5.27	4.93	1 47	2.35	10.70	87.07	2 50	2 6	3.07	1.38	4.03	2.05	2.85	5.00	69)	
6.82	6.80	6 70	9 6	6 . v	6/.9	6.77	6.77	6.74	6.72	6.72	6.71	6.66	6.65	6.65	8 64	j 6	0.0	6.62	6.62	6.59	6.57	6.54	6.53	6.57	1 1 1	0.00	9 6		9.0	0.40	6.45	6.43	6.43	6.42	6.41	6.41		r S	13
8.14	97.6	2.0	0.0	0.34	8.93	27.70	1.65	10.08	0:30	3.61	7.44	65.62	18 79	4.67	, i	67.7	4.52	0.07	63.60	0.07	0.12	0.42	7.31		4.04 4.04	0.24	6.55	8.71	8. 44. 7	0.12	7.26	27.63	1.70	0.45	6.74		0.07	0.21	
119	5 7	14.0	0.01	0.05	1.32	4.09	0.24	1.50	9.0	0.54	+	9.85	2 8 2		2 !	1.1/	0.68	0.0	9.61	0.01	0.02	90.0	1 13	4 3	0.41	9	1.01	7. ¥	1.30	0.05	1.13	4.30	0.26	20.0	. 4	9.5	0.13	0.03	
22020	K£3033	H56903	AA256231	H71230	H08595	AA010065	AA454562	AA454597	H17139	A A B C 0 7 2 3	07/50000	A 4 6 6 6 0 3	240000	AASSAOSA	AA456695	AA424790	H59203	AA448400	T55728	AA884622	A A D 3 Q 8 5 1	HOREAK	10001	12084/	N48075	T51617	R43328	AA953357	H21943	AA428186	AA045508	AA626788	AA034115	0.00000	74.07.04	AA448664	AA757564	H15910	
404046	131310	204580	681906	214604	46091	359119	809503	811582	54083	103488F	1031663	1,23011	0147410	950445	813149	768172	204214	781362	73638	1468597	375827	13667	43002	01400	281733	72441	32134	1573251	160233	773564	487297	877784	430007	10000	00400/	786078	395955	159462	
	413	4 4	415	416	417	418	1 017	2 5	3 5		77 6	57	474	425	426	427	428	429	430	73.	2 5	754	?	4 25	435	436	437	438	439	4 4 04	4	442		? :	‡	4 5	446	447	

Table 2-11: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Lines Genes upregulated 5 fold or more in at least 2 cancer cell lines.

8	7	7	7	7	7	7	7	2	2	2	7	7	7	7	7	73	7		, ,	4 (V (v	v (4 0	7 (۷ ،	v 0	, (7 (N (N (7 (2	~	8	
12.22	4.22	8.37	12.04	6.13	11.23	9.86	6.32	8.69	8.17	7.26	8.95	7.88	10.08	13.49	5.30	10.52	5 48		9 0	9 1	6.77	8.78	6.60	7.78	10.29	9.56	9.0g	4.83 0.1	7.48	7.07	5.03	7.86	7.19	5.37	5.40	
90.9	6.40	7.88	5.53	12.20	6.38	8.03	10.78	6.93	8.05	10.13	2.46	8.82	6.21	5.01	0.27	5.05	70.07	\$.2.0 \$.00 \$.00	8.92	7.60	0.76	5.59	, 10.69	2.18	7.55	2.69	7.76	5.90	9.73	8.86	11.73	6.77	4.55	8.09	3.03	
0.90	8.52	2.87	1.54	0.76	1.48	1.06	1.79	3.26	0.E0	1.61	7.40	2.11	2.51	76.0	12.18	2 6		0.87	0.68	1.35	11.05	4.19	1.20	8.48	0.57	6.15	1.59	7.62	1.12	2.39	1.52	3.55	6.43	4.66	9.62	
6:39	6.38	6.37	6.37	9.99 3.99	96. A	 3. 1.	e.3.1	9 6	6.29	6.29 6.28	0.40	6.27	7.0	4 6	0.43	0.43	C7'0	6.23	6.22	6.20	6.20	6.19	6.16	6.15	6.13	6.13	6.13	6.12	6.11	6.11	6.09	90.9	90.9	6.04	6.02	4
1.66	4 82	3.76		t 6	 	7 0	4.0.4 AO 4.0	51.30	5. 1. 5. 1.	, c	17.7	1.57	 0. 1.00	0.0	67.7	13.69	2.35	4.1	1.36	12.01	90.0	12.17	0.33	2.57	15.94	12.43	2.89	0.14	2.40	16.02	26.54	4.93	0.08	7.89	0.94	
0.26	22.0	0 0	0.0	5 6	0.24	5.	77.0	8.25	0.72	0.93	C:3	0.22	0.25	0.02	0.37	2.19	0.38	99.0	0.22	1.94	0.01	1.97	0.05	0.42	2.60	2.03	0.47	0.02	0.39	2.62	4.36	0.81	0.01	13	0.16	
. 8000	or second	AA431/53	N39611	AA405533	R93068	AA465223	AA459244	AA042812	N63744	W85878	N63988	N77828	H54659	R44201	N24869	AA430367	R11605	AA701978	R38539	H22568	W31389	AA447662	N56875	AA865924	AA630104	AA430744	AA609891	H18017	R92186	AA430052	AA459106	H96392	AA102068	AA425404	AA620437	
. 4444		782476	277112	772410	196824	814211	814444	486538	292936	416075	289496	302221	203179	34254	267135	769857	130004	435957	23073	52128	320602	813586	277476	1470195	854701	770992	1032080	50983	195817	781109	814303	256680	489805	773147	951068	
977	? :	449	420	451	452	453	454	455	456	457	458	459	460	461	462	463	464	465	466	467	468	469	470	471	472	473	474	475	476	477	478	479	480	7 4 5	482	

Table 2-11: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 5 fold or more in at least 2 cancer cell lines.

7	7	8	8	7	8	2	, ,	4 (N	7	7	8	7	7	8	7	•	י ו	v	7	7	7	7	7	84	8	7	8	2	7	8	7	7		4 (7 (7
5.20	7.25	7.87	7.79	6.52	. R.	60 8	1 6	Ø./0 1	7.53	2.86	8.22	8.94	7.60	6.64	69.6	8 71	1 1	90.7	6.22	5.62	5.26	8.09	6.29	5.11	1.55	8.40	7.45	5.77	10.01	5.36	4.86	1.55	9 54		/o.r	7.73	7.68
9.92	9.75	7.62	8.57	70	7 7	G. C	0.00	7.34	3.15	8.19	8.58	6.81	5.43	2.32	1.00	200	7	9.30	7.27	9.55	10.89	5.86	6.03	10.22	10.45	7.72	7.30	9.05	6.27	9.62	5,66	9.78	8 64	0.9	9.35	8.12	5.70
2.89	26.0	2 44	1 57		2.30) 6.0)	11.41	1.63	7.03	99.9	0.84	1.87	4.55	8.54	8.78) (6.50 0.50	4.53	3.92	2.24	1.21	3.39	4.97	1.96	5.27	1,10	2.46	2.42	68. 68. 68.	2.14	 	50.0 7.7	3 6	0. 0.	5.83	1.17	3.60
9	9 00	6 6 6 6 6 6	9 0	0 0 0	9. S	5.93	5.93	5.91	5.90	5.90	5.88	5.88	5.86	883) i	20.0	5.81	5.80	5.80	5.80	5.79	5.78	5.76	5.76	5.76	5.74	5 7.4		5.73	1 7	- 7.0	5 G	9.60	5.68	5.68	5.68	5.66
7. 7		9.00	φ (2.99	0.55	0.69	1.85	15.84	3.18	0.36	0.76	, r.) ()	7	50.4	90.00	43.39	10.85	32.21	4.83	0.22	13.63	0.16	0.59	4 73	2.62	i (5.37	9.16		0	د/.4 د ر	1.58	5.71	2.92	2.35	5.08
,	67.1	1.48	1.39	0:20	0.09	0.12	0.31	2.68	0.5 4	900	5 6		5 to	6.5	0.09	0.01	7.47	1.87	5.55	0.83	40.0	2.36	80	9 0	5 6	7 o	9 (D !). O. G	0.24	0.03	0.83	0.28	1.00	0.51	0.41	0.90
	AA458646	AA465353	AA504202	N39233	AA879423	AA101173	N62866	N36008	81992N	9180314	0000001	AA770003	N/3252	W84008	AA709271	H02158	AA454585	N59136	AA625662	AAARANGO	AA609914	T67360	10/333	TU1332	500001	N52128	K13434	R50752	H10788	H3/81/	R40481	AA150777	R45952	N72274	N51498	R89363	A A 228130
;	813410	814080	825284	276972	1505686	563860	289645	272888	271076	007100	204703	1040349	246549	35/531	1343468	150623	809535	287637	745360	814005	1031113	74738	44738	147.303	200403	28/581	284/5	38510	46862	191546	27817	504826	34063	291374	282104	196037	867608
	483	484	485	486	487	488	489	490	3 5		76	583	\$;	495	496	497	498	499	200	3 6	2 2	3 5	3 3	2 4 5	200	208	207	20g	503	510	511	512	513	514	515	516	£47

Table 2-11: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Linei Genes upregulated 5 fold or more in at least 2 cancer cell lines.

7	7	7	7	7	7	8	2			4 (7 (7	7	7	8	7	8	2	. ~		• (٠, ١	7 (7 (7	7 1	7 (7 '	N (7	74	7	7	7	7	7		
6.19	5.00	8.55	11.39	2.59	5.48	6.45	1 48	9 00	n (0. i	2.47	7.67	5.02	8.52	6.45	8.81	. 65.55	7 60	1.63		19.7	6.30	7.49	7.59	8. 44.	5.51	8.69	4.14	7.71	8.77	5.87	7.27	6.98	8.24	5.55	6.26		
2.36	1.37	77.7	97	8 42		2 0	o (07.1	10.00	9.88	7.41	2.85	6.93	7.04	20.8	000	S 6	87.0	47.7	40.	8.14	8.29	6.97	5.09	6.79	8.15	5.96	5.49	69.9	5.32	5.58	3.79	5.22	6.68	7.08		5	
8.40	10.55	, F.F.	, c	3.02 2.02	5.63	6.88	5.49	8.06	0.80	1.43	6.89	6.12	4.68	102	- c	2 [79.7	3.62	0.87	7.11	0.67	1.75	1.86	3.64	1.09	2.62	1.61	6.63	48.1	2.06	4.66	5.05	3.94	1 13	3	ה ני יי	5.00 00.5	
بر ور) (4)	* (0)	5.62	5.62	5.62	5.62	5.61	5.60	5.60	5.59	5.59	5.55	7 7	r c	0.00	5.52	5.49	5.49	5.49	5.48	5.47	5.45	5.44	5.44	5.44	5.43	5.42	5.42	5.41	5.38	5.37	5.37	72.37	, w) i	رن بن بن	5.33	16
10 63	6.6	6.68	0.25	0.10	90.0	25.20	5.38	17.75	0.07	0.17	4.03	3.65	9 6	0.40	132.57	0.41	66.94	5.08	11.11	8.93	5.38	0.26	0.05	0.31	1 05	0.45	33.51	5.65	0.59	7.13	, t		- 7	2.5	19.11	0.55	0.94	
9	3.28	1.19	0.05	0.05	0.01	4.49	96.0	3.17	0.01	0.03	22.0	99	0.00	0.07	23.98	0.07	12.19	0.93	2.02	1.63	0.98	0.05	50	900	3 6	e e	5 5 6	\$ 5		4 23		0, 0	2.10	08: ·	2.17	0.10	0.18	
	AA699560	AA669218	AA757588	AA427400	W93500	R80790	H93424	AA608730	14/96197	127246	14/0000H	1 96201	AA035580	H23983	AA670408	AA907714	AA487608	AA115559	R22308	H80685	D02812	AA428514	100250	000000	AA01/213	K8926/	AA884420	AA706629	AA596603	A4663100	AA45882/	AA779383	AA450205	AA682545	AA402965	AA676840	W93682	
	433474	856592	395609	616011	357344	146882	220700	950926	361688	20100	193139	98/171	471707	51465	878788	1505898	839081	491615	130845	241432	107206	13170	011011	220120	90105	195/86	1400990	401000	898097	140//01	814350	459947	789204	431245	741790	460114	357285	
	518	519	520	521	522	523	£2.4	£25	3 2	9 70	25/	228	223	230	531	532	533	253	535	838	2 2	200	22	539	54 04	1	542	2	4 :	<u>د</u> د	546	247	548	549	220	551	552	

Table 2-11: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes Upregulated 5 fold or more in at least 2 cancer cell lines.

8	73	7	7	8	7	7	7	7	8	7	7	8	5	7	7	7	7	. 6		• (4 (4 0	74 (7	2 9	71	7	8	74	Α.	7	7	73	7	8	
6:39	6.03	7.16	7.71	3.90	60.6	5.20	7.18	5.27	6.95	7.50	7.02	8.43	6.43	7.68	6.64	6.90	स्ट स्ट	6.97	, c	S	49.7	5.93	8.34	5.28	5.63	5.12	7.63	0.32	5.48	5.67	7.92	5.33	5.26	5.79	1.87	
5.23	1.01	7.63	7.20	5.91	5.86	9.61	6.76	8.00	6.68	2.95	2.97	5.07	7.32	5.49	5.99	6.91	***	u	9.5		0.80	5.12	5.30	8.40	5.03	9.05	5.29	8.82	8.12	8.65	5.51	6.66	3.76	8.28	6.10	
4.34	8.91	1.15	1.02	60.9	75	, c	60	2 49	2 22	5,35	5.79	2.21	1.93	2.51	2.95	1.73	: ·	1.23	3.25	7.12	6.70	4.34	1.68	1.62	4.62	1.14	2.35	6.08	1.54	0.82	1.63	3.05	6.00	0.91	7.00	
5.32	5.32	5.31	A 34) K	0.00 0.00	9.23 7.30	9 CC	5.78 8.78	5.26	5.26	5.24	5.23	K 23			<u>.</u>	5.17	5.16	5.15	5.15	5.13	5.11	5.10	5.09	5.09	5.09	5.07	5.05	5.04	5.02	5.02	5.01	4.99	4.99	17
35.80	0.05	6.28	0.15	1.7.	5.04	27.66	6.03 80.4 80.03	0.23	20.12	. 4. 1. 8.	9 5	† 0 - 4 - 6	34.05	5 4 4 3 5 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7	13.23	0.53	4.0	0.51	2.47	0.05	0.06	31.82	0.19	0.43	0.77	32.60	1.98	16.55	40.76	0.22	3.45	23.47	4.55	4.54	8.53	
6 73	5	- α - τ	0 (9.9	7:27	4.09	0.76	9 1	3.81	0.08	2 6	3 3	- 6 - 6	9 6	2.92	0.0	1.83	0.10	0.48	0.01	0.01	6.20	0.04	0.08	0.15	6.40	0.39	3.26	8.07	900	0.69	4.68	6	5 6	1.7	
A A & 20 O K K K	73575	K/3525	K166/6	AA598668	H98822	AA455652	N21334	N57858	177840	AA628190	H59305	AA879474	AA620614	AA133166	AA169645	AA446748	AA644334	AA426561	AA136060	AA398218	W84774	N66132	AA683338	AA777779	AA885851	AA148505	R71689	AA962541	AA421230	A 4 4 9 0 5 0 5	N49899	44504272	A 4 2 2 6 5 2	1646E3	AA149637	
7.7.7.7.7.7.7.7.7.7.7.7.7.7.7.7.7.7.7.	021108	1004/3	129664	898258	261714	813997	265102	247089	108395	1055831	204299	1505735	1048685	490753	594517	784126	845602	768146	490060	726658	415806	278483	1293016	448920	1500388	491524	155575	1602209	739247	824109	243653	825411	E64803	20000	50003) - -
c u	2 :	ž,	222	556	227	558	559	260	561	295	563	20	565	999	267	568	569	570	571	572	573	574	575	576	577	578	27.0	280	, g	283	4 6	200) u	000	200 7.87	}

Table 2-11: Prostate Cancer Cell Lines vs. Normal Prostatic Epithellaticell Lines appregulated 5 fold or more in at least 2 cancer cell lines.

8	7	64	7	7	7	2	7	6	. 6			4 (4 (7	7	7	7	7	7	7	^			4 6	v	7 (77 (N (2	7	7	64	7	7	8	8	ı	
6.61	4.03	5.79	0.97	5.65	5.61	6.75	6.79	700	333)))	0.60	04.0 04.0	٥٠./	5.90	6.59	1.00	5.64	5.32	1.00	6.60	7.30	03.7	2.02	5.63 5.63 6.63	6.25	6.47	6.37	8.48	6.21	6.12	5.01	5.80	4.	3.64	80 80 80 80 80 80 80 80 80 80 80 80 80 8) o	5.0	
6.33	5.47	8.07	5.74	8.14	5.83	9 6 6 8 8	o. 62	0.16	15.7	68.7	6.45	7.62	5.39	5.34	5.47	5.04	5.49	8.39	5.50	, k) () () ()	5.26	6.24	6.79	6.82	6.47	5.38	0.08	5.74	5.51	3.86	1.33	5.90	5.23	263	, i, i,	9.55	
1.99	5.36	0.98	8 10	0.97	3 37	3.32	- 6 - 6	02.1	5.15 	0.37	1.97	1.57	2.09	3.38	2.45	8.44	3.32	0.71	7 93	- 6. 6	7:17	1.94	2.51	1.60	1.18	1.30	2.49	5.66	2.27	2.52	5.15		6.62	90 50		Q (2.19	
4.97	4.96	4.95	90 7	t (5	76.4	28.4	4.91	4.90	4.90	4.90	4.89	4.88	4.88	4.87	4.84	4.83	4.81	4 80		08.4	9.4 O	4.80	4.79	4.77	4.75	4.75	4.75	4.74	4.74	4.72	4.67	4.67	4 65	 	0.4	4.65	4.63	18
1.93	0.75	3.4	5 6	0.00	8.73	11.53	0.30	2.11	0.08	8.32	0.71	1.49	34.71	2.66	10.43	0.05	08.0	13 00	5.33	0.05	92.90	7.55	24.62	3.44	1.68	0.60	4.36	16.44	0.31	4.54	3.48	5 C	2.00	7	99.	1.82	4.18	
0.39	0.15		07.1	0.0	1.78	2.34	90.0	0.43	0.02	1.70	0.14	0:30	7.11	0.54	2.16	5	5 6	2 6	18.7	0.01	19.35	1.57	5.14	0.72	0.35	0.13	0.92	3.47	0.06	96 0	0.00	t 8	50.0	0.00	0.36	0.39	0.90	
ONFOCIA	00000	AA626/84	K44346	H39560	AA478542	AA443969	R82041	N56982	AA114106	H16743	AA904969	R55750	A A N 3 5 3 8 4	AA418918	N62372	4 4 4 5 4 5 5 4	AA466600	AA156589	AA064627	R43271	AA456931	N31641	AA131526	R54846	H55915	AA521411	R00275	AA779165	AA700415	H06768	200700	028001	AA/0158/	AA676590	AA046321	H08796	AA443094	
264163	201102	19/1/8	34294	186132	784772	757206	147925	277545	563201	50114	1504152	40721	471598	767994	200560	00000	90000	202290	382451	32517	838568	258966	503715	154472	204129	826089	123255	453005	460553	43876	43820	66406	432/32	882488	376764	46097	809466	
0	0 0	589	290	591	265	593	594	595	596	297	598	989		3 8	5 6	700	200	\$	605	909	209	608	609	610	. 1		. E	5.5	2 7 7	2 4	9 10	719	818	619	620	621	622	

Table 2-11: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes Upregulated 5 fold or more in at least 2 cancer cell lines.

7 6	4 (7	7	8	7	7	7	7	7	7	7	8	7	8	8	7	8	8			٠ ٨		: 0	, ,	4 0	, ,			4 C	4 (7	7	7	7	7		
	0.4.0 0.4.0	6.25	5.98	5.05	7.00	2.28	5.85	5.23	5.23	7.00	5,54	5.34	5.12	5.56	6.95	5.02	5.57	3.12	α 1 · 1	0.70	1.7.1) &	0.00	0.01	0.70 1.00	9. I.	0.23 7.43	- t. c	0.50 0.50	5.23	5.54	5.26	5.86	0.44	6.47		
5.63	5.08	5.20	5.24	5.72	0.75	5.06	6.62	7 47	7 20	, r. 53	28.6	. K	30.80	2.55	. r	6.17	- o	0.90	2 6	60°C	5.25	6.24	20.02	5.71	5.63	2.84	0.22	5.60	0.31	5.46	5.14	6.04	5.25	6.96	5.51		
2.36	3.28	2.32	2.55	2.99	66 6) ; ;	; ; ; ; ;	- ·	5 G		3 7	1.4.5		 	9.30	77.1	07.7	0.60	5.20	1.56	6.45	1.21	2.31	1.72	0.87	5.24	7.74	2.19	6.57	2.44	2.38	1.74	1.92	5.57	66.0		
4.62	4.60	4.59	4.59	7.50	 	0	0.4. v	40.4	4.54	4.54	4 4		SC. 4	4.52	4.50	0.4	4.49	4.48	4.48	4.48	4.47	4.46	4.44	4.44	4.42	4.42	4.42	4.40	4.39	4.38	4.35	4.35	434	4 33	4.32	30° -	10 00
5.17	11.07	15.49	5 4 7 4 7 4) (4.50	0.06	12.03	13.59	5.05	2.67	0.38	2.96	13.59	40.40	2.71	6.39	4.53	1.60	10.51	0.32	1.09	0.53	90.0	0.49	1.88	0.08	7.57	57.53	0.14	0.84	0.73	0.40	0.40	0.42	7 6	 	
1.12	2.40	3 3 2	5.5) (a)	0.98	0.01	2.64	2.99	1.1	0.59	0.08	0.65	3.00	8.95	0.60	1.42	1.01	0.36	2.35	0.07	0.24	0.12	0.0	0.11	0.42	0.05	1.71	13.07	0.03	0.19	0.17		3 6	5 6	0.02	0.25	
AA626847	A A 4 72076	AA173920	AA598797	AA460521	AA284291	AA629251	AA676296	H98655	AA488979	N51367	AÁ704858	R24451	AA157787	AA421270	AA885096	H58542	T58434	N29624	AA454947	AA778562	H22854	AA883660	H20743	AA904776	R89765	R06372	AA398355	T90621	AA404338	04413070	A A 884382	70004307	AA62624/	AA625791	AA443300	AA205320	
745192	10101	801080	898083	796606	327245	744044	882355	261567	824728	283208	453229	132954	588840	731031	1468466	206272	75650	257312	814765	1048899	51700	1466599	51328	1504005	194811	126239	726858	110772	758302	ESOEAR	320343	01/10#1	745523	744911	784589	647816	
233	3 6	624	625	929	627	628	629	630	631	632	633	634	635	636	637	638	639	640	140	642	643	. 7	645	948	64. 54.	648	649	650	55		700	20 1	654	655	656	657	

N N	8	7	7	7	2	8	8	7	2	7	. 7	. 6	. 6	2	. 0	. 7	8	2		• (7 (2	
5.84 5.62	5.79	6.07	5.36	5.76	5.87	5.58	5.36	5.41	5.49	5,65			 	4 4 5	x	0.56		, , , , , , , , , , , , , , , , , , ,) u	66.0	5.01	5.63	
5.60	5.75	561	4,5	5.55	1 07	 	57.3	t) ·	6.03		5/.6	5.12	5.C	57.5	0	2.07 2.04	† (C)	5.12	5.69	5.60	5.56	
1.50	 	97:	9 6	- 4 - 4	C. 2. 7	0. 0	 5	7.58 8.51	0.50 0.50	9.84 4	1.58	0.57	1.23	0.97	5.88	5.25	5.37	1.13	1.12	5.46	1.01	0.43	2
4.32	4.28	4.28	4.25	4.24	9 9	4.17	4.16	4.16	4.14	4.13	4.12	4.10	4.09	4.08	4.08	4.01	3.92	3.91	3.90	3.90	3.87	2 87	ò.
22.74	0.39	0.27	6.07	11.37	8.82	5.73	11.43	0.36	1.69	9.76	0.57	9.78	0.83	4.55	4.25	7.30	0.07	10.81	2.97	0.09	1.56) - -	7.56
5.27	0.09	90.0	1.43	2.68	2.10	1.37	2.75	0.09	0.41	2.36	0.14	2.38	0.20	1.12	<u>5</u> .	1.82	0.05	2.77	0.76	0.02	9 6	?	1.95
T70122	AA437245	H08227	N50828	AA465692	N35579	R60949	N50935	AA883755	AA708431	AA487206	R07012	AA633993	AA862814	T54527	T62552	N79230	H79566	AA676598	H84915	NEB136	000000	H4/088	AA045340
80946	757463	45514	280909	814906	272238	42400	281103	1466824	505997	841287	126695	858292	1469211	73659	79726	292388	239661	882483	249517	247597	100147	193420	487141
658	629	099	199	299	663	664	665	999	299	668	699	670	67.5	672	673	674	675	676	677		9/9	679	680

Table 2-12: Prostate Cancer Cell Lines vs. Normal Prostatic Epithellal Cell Line. Genes upregulated 10 fold or more in at least 2 cancer cell lines.

Table 2-12: Prostate Cancer Cell Lines vs. Normal Prostastic Epithelial Cell Line: Genes upregulated 10 fold or more in at least 2 cancer cell lines.

Order	IMAGE	Gen Bank Accession	MPM1(PrEC)	Ave Cancer Cell Lines	Fold (Ave Cell line/MPM1)	Fold(MPM2/M PM1)	Fold(MPM271/M M1)	Fold(MPM2/M Fold(MPM271/MP Fold(MPM272/M # of Cancer PM1) PM1) Cell Lines	# of Cancer Cell Lines
	Clone IO	Number		Values		13 45	301.13	22.85	ო
-	1343732	AA725564	0.01	1.12	112.48		106.60	43.45	ო
8	784296	AA447079	0.01	0.66	65.30	,	138.91	35.70	ო
က	796569	AA460463	0.01	0.35	63.5/	02.00	17.92	47.97	ო
4	1558655	AA976561	0.08	4.36	54.32	9 4 4 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5	123.80	11.93	ო
2	743016	AA406036	0.01	0.50	50.47	0.00	20 10	50.51	က
9	33076	R44048	0.80	36.21	45.35	20.30	20.62	22.90	က
7	626390	AA189106	0.07	3.16	45.16	20.18	80.79	14.51	ю
ω	812170	AA456035	0.01	0.30	37.18	16.25	87.45	10.68	က
Ø	50288	H17888	0.01	0.32	36.79	12.25	30.44	28.82	က
5	487861	AA045436	0.02	0.59	32.57	29.44	38.30	24.80	ო
=	291057	N72115	0.23	6.85	29.19	24.40	2.50	41.29	ო
12	263014	H99813	0.63	17.34	27.31	19.07	26.0 <u>2</u>	31.77	ო
13	303180	N92764	0.01	0.27	27.21	13.82	20.00	13.87	ო
4	46896	H09818	0.05	1.21	25.96	33.86	30.10	25.64	ო
5	151184	H02294	0.48	12.53	25.93	38.89	13.20	12.48	ю
16	41595	R59556	0.01	0.17	25.59	12.06	12.60	26.60	ო
17	430092	AA009840	0.40	10.00	24.80	34.18		10.88	ო
⊕	1049033	AA778675	0.38	9.30	24.51	46.38	10.20	32.50	ო
19	884790	AA629838	0.01	0.24	24.29	30.26	16.09	11.55	ო
20	365149	AA025142	0.16	3.52	22.24	38.20	10.30	24.62	ო
. 72	51581	H22824	0.01	0.25	20.42	18.18	16.45 24.35	20.16	က
22	1467161	AA883187	0.02	0.31	18.24	10.20	28.33	14.13	ო
ន	488431	AA047441	0.17	2.95	17.83	10.43	14.07	12.25	ო
24	82869	T69270	0.02	0.32	15.94	27.48	. 4. 5. 5. 5. 5. 5. 5. 5. 5. 5. 5. 5. 5. 5.	15.62	ო
52	898286	AA598974	1.21	16.52	13.65	13.47	64.14	1917.92	84
78	42627	R60995	0.01	9.14	660.46	2.32	66.27	1701.31	7
27	757165	AA443950	0.01	5.71	591.67	7.33	99.56	14 13	74
28	592707	AA160606	0.01	1.66	234.30	01.8 	674.66	35	74
53	781036	AA446446	0.01	1.70	228.86 1	10.58	. 00.4/0	<u>.</u>	

Table 2-12: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line Genes upregulated 10 fold or more in at least 2 cancer cell lines.

2	7	8		7	73	7	81	0	7	0	7	7	8	7	7	8	8	7	7	7	7	7	7	N	7	7	7	7	7	7	7	8	8	7	
10.82	146.14	105.59	12.54	13.76	99.88	11.76	70.45	123.70	0.72	12.17	5.49	91.51	6.93	11.05	19.33	12.68	51.22	9.41	11.01	55.73	48.14	43.19	14.72	11.03	11.15	52.45	36.33	63.66	14.11	10.31	16.63	11.62	31.35	45.50	
570.75	140.40	150.46	239.88	229.57	95.72	168.10	4.69	51.92	88.50	139.36	44.77	56.58	101.07	109.53	99.20	88.28	52.03	73.72	82.72	34.54	30.24	36.47	65.78	70.25	69.04	27.83	35.03	11.75	58.23	66.04	54.64	60.05	41.17	26.82	
8.10	0.80	0.89	3.09	1.42	0.87	3.73	104.24	1.03	78.84	0.63	99.52	0.91	25.03	4.59	4.81	5.91	0.93	20.19	5.81	0.77	9.10	7.73	5.99	0.74	1.21	69.0	7.36	2.28	5.18	0.59	4.46	3.61	2.32	2.27	
196.56	95.78	85.65	85.17	81.58	65.49	61.20	59.79	58.88	56.02	50.72	49.93	49.67	44.34	41.72	41.11	35.62	34.73	34.44	33.18	30.35	29.16	29.13	28.83	27.34	27.13	26.99	26.24	25.90	25.84	25.65	25.24	25.09	24.95	24.86	2
1.28	193	6.83	0.51	0.82	5.16	0.61	26.93	2.87	12.41	1.10	5.02	13.02	0.44	0.42	0.41	0.36	8.56	0.25	0.33	7.89	0.40	0.41	0.44	0.27	0.27	14.38	0.16	14.15	0.24	0.39	0.25	0.17	0.25	24.91	
0 0	200	800	600	5 6	800	9 6	0.45	0.05	0.22	0.02	0.10	0.26	0.01	0.01	0.01	0.01	0.25	0.01	0.01	0.26	0.01	0.01	0.02	0.01	0.01	0.53	0.0	0.55	0.01	0.02	0.01	0.01	500	9:	
702693	LEGARA	004000	AA450414	M4439917	V60011	W4 19400	D23270	AA251784	W73144	R44741	R51617	AA504844	R09747	AA437126	AA810225	H94482	R91566	8600EN	AA608832	AA485052	AA863292	77779	AA789328	R15740	AA460274	H50655	AA418903	N47445	R49555	N52767	AA010383	AA236986	WOAABE	H19826	
220064	220033	845704	705627	347182	746462	746152	131452	684879	344589	33827	39442	825740	128083	757368	1367900	243181	196544	256911	1030613	815861	1455835	108378	1391682	53039	796549	194318	768056	280699	38344	283379	430255	687972	357884	172517	
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11.25	43.13	9.98	48.25	26.43	25.34	31.76	11.14	54.20	23.77	3.08	47.63	22.42	17.48	9.30	28.27	20.17	45.41	40.27	10. 10. 10. 10. 10. 10. 10. 10. 10. 10.	1.08	80:- 90:-	15.05	16.77	10.22	, 4 , 8 8 8	12.21	38.04	31.84	34.18	33.48	26.82	28.36	25.45	10.72	9:64	
60.48	28.65	17.55	21.91	24.50	42.67	30.37	5195	Se C	10.03	5 6	23.72	10.00	100 100 100 100 100 100 100 100 100 100	30.57	20.00	38 73	200.	10:01	11.10	30.81	14.51	70.12	38.83	34.10 0.00	33.82	- 8.7° - 8.7° - 8.7°	44.78	10.67		10:1	9.03		16.92	80.12 85.69	77.79	
1.76	0.95	44.37	5.50		- 6 - •	90:- 7	9 0	3.82	0.88	20.95	37.52	4.40	o	0.93	20.94	87.L	00.1	2.79	7.04	2.35	41.41	3.39	4.49	1.97	7.52	25.95	1.87	49.6	0.70 7.8	0.4	10.47	5.81	2.63	0.69	4.69	
24.50	70.70	+7:+7	23.97	23.68	23.01	23.00	22.37	22.30	21.71	21.70	21.44	20.97	20.40	20.39	20.27	20.04	19.97	19.50	19.50	19.47	18.93	18.85	18.01	17.44	17.36	17.25	17.19	17.15	16.89	16.78	16.35	16.17	15.84	15.64	15.57	ო
700	\$7.0 7.1	7.41	0.16	3.51	0.16	0.23	0.16	3.74	3.65	0.14	40.77	19.16	0.20	6.19	2.46	3.94	0.20	0.20	0.19	3.30	21.01	0.17	0.35	0.17	0.21	33.78	1.06	0.63	0.17	0.11	0.16	11.31	2.94	5.98	0:30	
	D.O.	0.31	0.01	0.15	0.01	0.01	0.01	0.17	0.17	0.01	1.90	0.91	0.01	0.30	0.12	0.20	0.01	0.01	0.01	0.17	1.1	0.01	0.02	0.01	0.01	1.96	90.0	0.04	0.01	0.01	0.01	0.70	0.19	0.38	0.02	
	H96647	AA490945	R39191	AA504625	N63057	T70198	AA630328	R44936	AA491261	H10072	T61938	N32904	N32542	AA255954	H22563	R95684	AA461508	R86970	R54036	H28091	T65736	R54492	AA057195	AA001376	AA701315	T49355	AA857163	N47012	AA676998	AA481481	W92263	W93717	AA504838	AA417713	AA045175	
	251806	824025	23345	825606	284714	81050	855910	34204	824270	46843	79000	259884	270975	686552	51916	199337	795831	197265	40010	162310	80338	39722	489076	361899	435099	67440	1410444	280155	454083	756629	358990	357373	825726	746232	487766	
	65	99	29	68	69	20	71	72	5	74	75	292	11	78	79	80	81	82	83	3		98	87	88	8	06	91	92	83	8	95	96	26	86	66	

Table 2-12: Prostate Cancer Cell Lines vs. Normal Phostatic Epithelial Oell Line. Genes upregulated 10 fold or more in at least 2 cancer cell lines.

811766 51599 250069	AA463445 H18932 H97140	0.26 0.26 0.02	3.94 0.35 0.35	15.03 14.95 14.92 14.61	1.53 0.42 5.14 5.11	28.43 26.10 24.61 16.46	15.13 18.34 15.00 22.27	0 0 0 0 0 C
28298 785293 770860 1461609	K40434 AA476543 AA434388 AA883800	0.11 0.01 0.25	1.64 0.14 3.51	14.41 14.40 14.24	1.00	3.02 18.93 4.19	15.54 23.25 28.45 27.66	
418004 41406 1035796	W90705 R56149 AA628867	0.30	4.17 0.13 15.63	13.89 13.76 13.71	2.99 17.85 10.64	11.04 18.97 9.76 26.44	20.74 20.74 12.18	
26806 502003 771004	R37738 AA128617 AA427719	0.10 0.01 0.29	1.31 0.17 3.95	13.58 13.58 13.50	2.42 7.33 0.64 8.23	15.99 21.58 12.04	17.41 18.27 19.45	n n n
415089 824074 51103	W93379 AA491227 H19217	0.65 0.02 0.02	8.63 0.26 0.23	13.22	3.39	18.08 17.85 1.54	18.18 6.86 26.50	ппп
284288 361122 244050	N52193 AA017379 N34042	0.0	0.08 0.13 0.17	12.88 12.88 12.88	0.65 5.65 27.57	12.28 21.72 14.55	25.70 11.26 0.58	ппп
795446 1049185 364352	AA453616 AA620697 AA022496	0.01 0.76 0.01	9.44 0.12	12.39	1.96	0.39 14.19 23.87	26.39 20.98 11.09	ппп
47264 686172 1049079	H10713 AA262211 AA778717	0.30 1.18 0.03	3.72 14.39 0.31	12.18 12.10 12.00	5.65 3.28 1.00	13.80 15.47 14.95	17.09 17.55 20.10	000
195845 451649 739450	R92201 AA706901 AA477227	0.14 1.99	1.69 16.82 23.53	11.98	2.08	18.70 17.13	12.88 16.24 19.11	0 0 0
46461 1030798 591157	H09940 AA609002 AA161188	0.33 0.03 0.02	3.89 0.31 0.18	11.80 11.73 11.71	10.15 0.58	8.33	16.72	0 0 0
33839 357138	R44816 W93523	0.96 0.02 2.38	11.24 0.20 27.64	11.67 11.63 11.61	3.60 10.87 3.65	19.51 20.61 10.18	3.40	1 61 61 6
592728	AA160670	0.03	0.34	11.17	0.33	16.85	÷>:0:	ı

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13.80	16.62	10.20	12.02	20.10	12.92	7.26	15.60	15.64	1.65	17.42	16.30	14.06	15.31	10.88	7.37	10.89	5.79	11.50	15.44	16.07	15.05	44 44	5.24	4.33	130	13.51	13.08	14.35	14.67	1.10	16.60	1.94	14.35	11.27	<u>:</u>	
17.96	15.37	20.21	15.78	10.22	8 10	13.18	14.48	14.23	17.15	11.85	12.15	3.07	13.69	17.62	10.28	15.63	20.07	11.73	1.1.1	69.1.	1.00		12.53	16.44	5 7	50.11	10.73) ye	10.87	13.45) (2)	96. 96.	56.7	5.7	04:71	
0.92	0.51	50.6		; c	11.07	0:- 7	9.0	4 8	7.7	10.11	6 ×	42.44	16.7	5 •	5 5	t ::	co.7	39.11	0.00	1.15	10.45	11.43	0.97	10.06	12.49	4.58	9.61	0+.6	12.50	3 . 5	12.04	0.00	27.11	10.33	1.45	
10 90	, c	t 60.00	0.03	10.81	8/:0L	10.70	10.64	10.32	10.21	10.11	70.0L	C 10.00	6.60 6.60	D 1	9.87	9.57	9.53	9.52	9.48	9.48	9.47	9.34	9.31	9.24	9.22	9.17	9.12	60.6	9.07	9.6	9.03	8.98	8.87	8.56	8.39	လ
c c	5.80 0.00	51.9	0.31	90.0	9.32	60.0	13.51	0.87	5.55	0.12	2.53	2.29	1.05	4.94	3.82	0.16	0.77	10.61	1.40	1.40	0.09	90.0	0.22	1.03	0.97	9.89	2.44	0.26	0.26	3.21	11.71	17.81	5.97	0.09	9.51	
6	0.36	0.57	0.03	0.01	0.86	0.01	1.27	90.0	0.54	0.01	0.25	0.23	0.11	0.50	0.39	0.02	90.0	1.12	0.15	0.15	0.01	0.01	0.02	0.11	0.10	1.08	0.27	0.03	0.03	0.35	1.30	1.98	29.0	0.01	1.13	
	H48070	AA410434	N74131	AA461513	N30751	AA453433	H58234	AA281784	N49389	AA158035	R94504	AA664007	AA625979	AA292226	N21633	AA701652	R46000	N47113	00779	AA219060	W16659	AA284072	N50962	H04789	AA705047	AA446462	R96579	R69584	W44766	N26515	AA172056	AA620485	W68220	AA496998	W47364	
	193333	753428	298417	795840	257955	788209	204483	712401	277621	590640	197651	855391	745490	725877	266135	433567	35804	280375	303089	630013	320495	700792	281162	43733	461488	781047	199577	141726	320865	266259	594758	951241	342640	897546	324717	
	135	136	137	138	139	140	141	142	143	1 4	145	146	147	148	149	150	151	152	153	15.4	155	156	157	158	159	<u>8</u>	161	162	163	2	165	166	167	168	169	

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	1.30	2.23	/0.rr
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	AA504265	AA058576	AA706968
	825416	380884	451907

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Comuence #	Accession #	Database	Sequence 52	AB033048	GENBANK
Sequence #	A21185	GENBANK	Sequence 53	AB033114	GENBANK
Sequence 1 Sequence 2	AB000509	GENBANK	Sequence 54	AD001528	GENBANK
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Sequence 4	AB002366	GENBANK	Sequence 57	AF002715	GENBANK
Sequence 5		GENBANK	Sequence 58	AF003837	GENBANK
Sequence 6	AB002387		Sequence 59	AF004561	GENBANK
Sequence 7	AB004304	GENBANK	•	AF006085	GENBANK
Sequence 8	AB004788	GENBANK	Sequence 60	AF007791	GENBANK
Sequence 9	AB004903	GENBANK	Sequence 61	AF007791	GENBANK
Sequence 10	AB007870	GENBANK	Sequence 62 Sequence 63	AF010235	GENBANK
Sequence 11	AB007899	GENBANK	Sequence 64	AF013988	GENBANK
Sequence 12	AB007930	GENBANK	Sequence 65	AF016266	GENBANK
Sequence 13	AB007941	GENBANK	•	AF016270	GENBANK
Sequence 14	AB007956	GENBANK	Sequence 66	AF010270	GENBANK
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Sequence 18	AB011108	GENBANK	Sequence 70	AF025998	GENBANK
Sequence 19	AB011132	GENBANK	Sequence 71	AF026939 AF027824	GENBANK
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Sequence 26	AB012910	GENBANK	Sequence 78	AF035309	GENBANK
Sequence 27	AB014521	GENBANK	Sequence 79	AF038404	GENBANK
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Sequence 49	AB029000	GENBANK	Sequence 101	AF057160	GENBANK
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Sequence 51	AB033020	GENBANK	Sequence 103	AF061938	GENBANK

Page 1 (of 73 pages in Table 3-1)

Sequence 104	AF067656	GENBANK	Sequence 156	AF155832	GENBANK
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Sequence 111	AF070655	GENBANK	Sequence 163	AJ004913	GENBANK
Sequence 112	AF070657	GENBANK	Sequence 164	AJ004955	GENBANK
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Sequence 114	AF071202	GENBANK	Sequence 166	AJ006834	GENBANK
Sequence 115	AF071593	GENBANK	Sequence 167	AJ007398	GENBANK
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Sequence 123	AF086218	GENBANK	Sequence 175	AJ238095	GENBANK
Sequence 124	AF086336	GENBANK	Sequence 176	AJ250042	GENBANK
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Sequence 128	AF086557	GENBANK	Sequence 180	AL049799	GENBANK
Sequence 129	AF087020	GENBANK	Sequence 181	AL049932	GENBANK
Sequence 130	AF087573	GENBANK	Sequence 182	AL049941	GENBANK
Sequence 131	AF088055	GENBANK	Sequence 183	AL049942	GENBANK
Sequence 132	AF091089	GENBANK	Sequence 184	AL049969	GENBANK
Sequence 133	AF093118	GENBANK	Sequence 185	AL049987	GENBANK
Sequence 134	AF100615	GENBANK	Sequence 186	AL050041	GENBANK
Sequence 135	AF100755	GENBANK	Sequence 187	AL050141	GENBANK
Sequence 136	AF100756	GENBANK	Sequence 188	AL050161	GENBANK ·
Sequence 137	AF101051	GENBANK	Sequence 189	AL050184	GENBANK
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Sequence 208	AL133000	GENBANK	Sequence 260	J03209	GENBANK
Sequence 209	D00015	GENBANK	Sequence 261	J03250	GENBANK
Sequence 210	D00017	GENBANK	Sequence 262	J03464	GENBANK
Sequence 211	D13641	GENBANK	Sequence 263	J03779	GENBANK
Sequence 212	D14041	GENBANK	Sequence 264	J03799	GENBANK
Sequence 213	D14696	GENBANK	Sequence 265	J04164	GENBANK
Sequence 214	D16431	GENBANK	Sequence 266	J04621	GENBANK
Sequence 215	D25215	GENBANK	Sequence 267	J04970	GENBANK
Sequence 216	D26488	GENBANK	Sequence 268	J05021	GENBANK
Sequence 217	D28358	GENBANK	Sequence 269	J05032	GENBANK
Sequence 218	D28420	GENBANK	Sequence 270	K01911	GENBANK
Sequence 219	D28482	GENBANK	Sequence 271	K02765	GENBANK
Sequence 220	D29954	GENBANK	Sequence 272	L00160	GENBANK
Sequence 221	D29958	GENBANK	Sequence 273	L01439	GENBANK
Sequence 222	D30037	GENBANK	Sequence 274	L02426	GENBANK
Sequence 223	D31839	GENBANK	Sequence 275	L05425	GENBANK
Sequence 224	D37965	GENBANK	Sequence 276	L06133	GENBANK
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Sequence 228	D43950	GENBANK	Sequence 280	L11066	GENBANK
Sequence 229	D50371	GENBANK	Sequence 281	L11932	GENBANK
Sequence 230	D63486	GENBANK	Sequence 282	L13210	GENBANK
Sequence 231	D63874	GENBANK	Sequence 283	L13385	GENBANK
Sequence 232	D80009	GENBANK	Sequence 284	L13773	GENBANK
Sequence 233	D83032	GENBANK	Sequence 285	L13799	GENBANK
Sequence 234	D83197	GENBANK	Sequence 286	L15702	GENBANK
Sequence 235	D83485	GENBANK	Sequence 287	L19161	GENBANK
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Sequence 237	D86228	GENBANK	Sequence 289	L19872	GENBANK
Sequence 238	D86322	GENBANK	Sequence 290	L20941	GENBANK
Sequence 239	D87437	GENBANK	Sequence 291	L22569	GENBANK
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Sequence 243	D87667	GENBANK	Sequence 295	L38941	GENBANK
Sequence 244	D87685	GENBANK	Sequence 296	L38961	GENBANK
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Sequence 246	D89053	GENBANK	Sequence 298	L48984	GENBANK
Sequence 247	D89667	GENBANK	Sequence 299	M10119	GENBANK
Sequence 248	D89937	GENBANK	Sequence 300	M11119	GENBANK
Sequence 249	D90359	GENBANK	Sequence 301	M12670	GENBANK
Sequence 250	D90427	GENBANK	Sequence 302	M13231	GENBANK
Sequence 251	E01497	GENBANK	Sequence 303	M13692	GENBANK
Sequence 252	E01956	GENBANK	Sequence 304	M15661	GENBANK
Sequence 253	E01979	GENBANK	Sequence 305	M15885	GENBANK
Sequence 254	E02628	GENBANK	Sequence 306	M15887	GENBANK
Sequence 255	E02822	GENBANK	Sequence 307	M16247	GENBANK
Sequence 256	E02823	GENBANK	Sequence 308	M16342	GENBANK
Sequence 257	E06721	GENBANK	Sequence 309	M16660	GENBANK
Sequence 258	J00194	GENBANK	Sequence 310	M16768	GENBANK
Sequence 259	J03040	GENBANK	Sequence 311	M17324	GENBANK
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			0 004	1100040	GENBANK
Sequence 312	M17517	GENBANK	Sequence 364	U09848	
Sequence 313	M17885	GENBANK	Sequence 365	U10439	GENBANK
Sequence 314	M18366	GENBANK	Sequence 366	U13616	GENBANK
Sequence 315	M21302	GENBANK	Sequence 367	U14394	GENBANK
Sequence 316	M22382	GENBANK	Sequence 368	U14510	GENBANK
Sequence 317	M22918	GENBANK	Sequence 369	U16799	GENBANK
Sequence 318	M22920	GENBANK	Sequence 370	U19557	GENBANK
Sequence 319	M24194	GENBANK	Sequence 371	U24105	GENBANK
Sequence 320	M24594	GENBANK	Sequence 372	U24704	GENBANK
Sequence 321	M24902	GENBANK	Sequence 373	U28964	GENBANK
Sequence 322	M25246	GENBANK	Sequence 374	U30521	GENBANK
Sequence 323	M26663	GENBANK	Sequence 375	U34252	GENBANK
Sequence 324	M27332	GENBANK	Sequence 376	U37143	GENBANK
Sequence 325	M27334	GENBANK	Sequence 377	U37283	GENBANK
Sequence 326	M27826	GENBANK	Sequence 378	U37519	GENBANK
Sequence 327	M28211	GENBANK	Sequence 379	U38654	GENBANK
Sequence 328	M28372	GENBANK	Sequence 380	U39360	GENBANK
Sequence 329	M29064	GENBANK	Sequence 381	U39840	GENBANK
Sequence 330	M29550	GENBANK	Sequence 382	U40671	GENBANK
Sequence 331	M29870	GENBANK	Sequence 383	U41515	GENBANK
Sequence 332	M34840	GENBANK	Sequence 384	U42594	GENBANK
Sequence 333	M36341	GENBANK	Sequence 385	U51134	GENBANK
Sequence 334	M37583	GENBANK	Sequence 386	U54558	GENBANK
Sequence 335	M55409	GENBANK	Sequence 387	U54831	GENBANK
Sequence 336	M55542	GENBANK	Sequence 388	U56255	GENBANK
Sequence 337	M55543	GENBANK	Sequence 389	U58855	GENBANK
Sequence 338	M58485	GENBANK	Sequence 390	U60975	GENBANK
Sequence 339	M59305	GENBANK	Sequence 391	U62961	GENBANK
Sequence 340	M62840	GENBANK	Sequence 392	U63810	GENBANK
Sequence 341	M63838	GENBANK	Sequence 393	U65928	GENBANK
Sequence 342	M69181	GENBANK	Sequence 394	U66616	GENBANK
Sequence 343	M74509	GENBANK	Sequence 395	U70735	GENBANK
Sequence 344	M74558	GENBANK	Sequence 396	U80034	GENBANK
Sequence 345	M74777	GENBANK	Sequence 397	U80743	GENBANK
Sequence 346	M76180	GENBANK	Sequence 398	U81001	GENBANK
Sequence 347	M77830	GENBANK	Sequence 399	U90028	GENBANK
Sequence 348	M80902	GENBANK	Sequence 400	U90144	GENBANK
Sequence 349	M83653	GENBANK	Sequence 401	U90916	GENBANK
Sequence 350	M83738	GENBANK	Sequence 402	V00478	GENBANK
Sequence 351	M83941	GENBANK	Sequence 403	X01742	GENBANK
Sequence 352	M85164	GENBANK	Sequence 404	X02422	GENBANK
Sequence 353	M87339	GENBANK	Sequence 405	X02457	GENBANK
Sequence 354	M92439	GENBANK	Sequence 406	X02544	GENBANK
Sequence 355	M94654	GENBANK	Sequence 407	X04098	GENBANK
Sequence 356	S70154	GENBANK	Sequence 408	X04412	GENBANK
v .	S71513	GENBANK	Sequence 409	X04470	GENBANK
Sequence 357	S82240	GENBANK	Sequence 410	X04503	GENBANK
Sequence 358			Sequence 411	X05276	GENBANK
Sequence 359	U02556	GENBANK	Sequence 411	X05332	GENBANK
Sequence 360	U03271	GENBANK	•	X05908	GENBANK
Sequence 361	U07681	GENBANK	Sequence 413	X05906 X07695	GENBANK
Sequence 362	U08815	GENBANK	Sequence 414		
Sequence 363	U09564	GENBANK	Sequence 415	X07820	GENBANK

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0	V40507	OTABANIZ	Seguence 468	Z74615	GENBANK
Sequence 416	X12597	GENBANK	Sequence 468 Sequence 469	AA005035	dbEST
Sequence 417	X13425	GENBANK	•	AA007280	dbEST
Sequence 418	X14420	GENBANK	Sequence 470	AA010019	dbEST
Sequence 419	X15187	GENBANK	Sequence 471	AA010019	dbEST
Sequence 420	X15729	GENBANK	Sequence 472	AA010003 AA012927	dbEST
Sequence 421	X15822	GENBANK	Sequence 473	AA012927 AA016274	dbEST
Sequence 422	X15880	GENBANK	Sequence 474	AA010274 AA018892	dbEST
Sequence 423	X16940	GENBANK	Sequence 475	AA010092 AA019498	dbEST
Sequence 424	X52022	GENBANK	Sequence 476		dbEST
Sequence 425	X53280	GENBANK	Sequence 477	AA029415	
Sequence 426	X54941	GENBANK	Sequence 478	AA031483	dbEST
Sequence 427	X55525	GENBANK	Sequence 479	AA031762	dbEST
Sequence 428	X57810	GENBANK	Sequence 480	AA033869	dbEST
Sequence 429	X59618	GENBANK	Sequence 481	AA034226	dbEST
Sequence 430	X61970	GENBANK	Sequence 482	AA034237	dbEST
Sequence 431	X62744	GENBANK	Sequence 483	AA035742	dbEST
Sequence 432	X63432	GENBANK	Sequence 484	AA037126	dbEST
Sequence 433	X64875	GENBANK	Sequence 485	AA037172	dbEST
Sequence 434	X65550	GENBANK	Sequence 486	AA037554	dbEST
Sequence 435	X66276	GENBANK	Sequence 487	AA039354	dbEST
Sequence 436	X70326	GENBANK	Sequence 488	AA039504	dbEST
Sequence 437	X71087	GENBANK	Sequence 489	AA040209	dbEST
Sequence 438	X73114	GENBANK	Sequence 490	AA041494	dbEST
Sequence 439	X73902	GENBANK	Sequence 491	AA043431	dbEST
Sequence 440	X74801	GENBANK	Sequence 492	AA044187	dbEST
Sequence 441	X75593	GENBANK	Sequence 493	AA044209	dbEST
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Sequence 443	X76180	GENBANK	Sequence 495	AA044691	dbEST
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Sequence 445	X85372	GENBANK	Sequence 497	AA045584	dbEST
Sequence 446	X87241	GENBANK	Sequence 498	AA047213	dbEST
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Sequence 459	Y08915	GENBANK	Sequence 511	AA076171	dbEST
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Sequence 464	Z23064	GENBANK	Sequence 516	AA081800	dbEST
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Sequence 466	Z29330	GENBANK	Sequence 518	AA082437	dbEST
Sequence 467	Z36785	GENBANK	Sequence 519	AA083654	dbEST
Joquenies 701	200100	OFIND/MIK	254431100 0 10		

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Sequence 520	AA086007	dbEST	Sequence 572	AA186733	dbEST
Sequence 521	AA088322	dbEST	Sequence 573	AA187659	dbEST
Sequence 522	AA088850	dbEST	Sequence 574	AA187817	dbEST
Sequence 523	AA092224	dbEST	Sequence 575	AA188918	dbEST
Sequence 524	AA099952	dbEST	Sequence 576	AA190676	dbEST
Sequence 525	AA100987	dbEST	Sequence 577	AA194988	dbEST
Sequence 526	AA101208	dbEST	Sequence 578	AA195560	dbEST
Sequence 527	AA113403	dbEST	Sequence 579	AA196978	dbEST
Sequence 528	AA115604	dbEST	Sequence 580	AA203237	dbEST
Sequence 529	AA120818	dbEST	Sequence 581	AA205870	dbEST
Sequence 530	AA121471	dbEST	Sequence 582	AA211498	dbEST
Sequence 531	AA121923	dbEST	Sequence 583	AA213548	dbEST
Sequence 532	AA125809	dbEST	Sequence 584	AA215800	dbEST
Sequence 533	AA125989	dbEST	Sequence 585	AA218858	dbEST
Sequence 534	AA126128	dbEST	Sequence 586	AA223425	dbEST
Sequence 535	AA127674	dbEST	Sequence 587	AA224230	dbEST
Sequence 536	AA128306	dbEST	Sequence 588	AA225025	dbEST
Sequence 537	AA128391	dbEST	Sequence 589	AA229993	dbEST
Sequence 538	AA129501	dbEST	Sequence 590	AA232738	dbEST
Sequence 539	AA129909	dbEST	Sequence 591	AA233867	dbEST
Sequence 540	AA132798	dbEST	Sequence 592	AA234149	dbEST
Sequence 541	AA132956	dbEST	Sequence 593	AA234464	dbEST
Sequence 542	AA133382	dbEST	Sequence 594	AA234646	dbEST
Sequence 543	AA133503	dbEST	Sequence 595	AA234667	dbEST
Sequence 544	AA134032	dbEST	Sequence 596	AA247638	dbEST
Sequence 545	AA134778	dbEST	Sequence 597	AA250846	dbEST
Sequence 546	AA135049	dbEST	Sequence 598	AA250854	dbEST
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Sequence 548	AA136794	dbEST	Sequence 600	AA255577	dbEST
Sequence 549	AA136830	dbEST	Sequence 601	AA256290	dbEST
Sequence 550	AA148002	dbEST	Sequence 602	AA256297	dbEST
Sequence 551	AA149624	dbEST	Sequence 603	AA256591	dbEST
Sequence 551	AA150928	dbEST	Sequence 604	AA258003	dbEST
Sequence 553	AA156040	dbEST	Sequence 605	AA261968	dbEST
•	AA159847	dbEST	Sequence 606	AA262683	dbEST
Sequence 554			•	AA262783	dbEST
Sequence 555	AA160002 AA161003	dbEST dbEST	Sequence 607	AA202763 AA278357	dbEST
Sequence 556			Sequence 608	AA278785	dbEST
Sequence 557	AA164473	dbEST	Sequence 609		
Sequence 558	AA165243	dbEST	Sequence 610	AA278956	dbEST dbEST
Sequence 559	AA165632	dbEST	Sequence 611	AA279145	
Sequence 560	AA165639	dbEST	Sequence 612	AA279262	dbEST
Sequence 561	AA166907	dbEST	Sequence 613	AA282956	dbEST
Sequence 562	AA169276	dbEST	Sequence 614	AA283112	dbEST
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Sequence 564	AA171516	dbEST	Sequence 616	AA291436	dbEST
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Sequence 567	AA180270	dbEST	Sequence 619	AA293027	dbEST
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Sequence 569	AA181580	dbEST	Sequence 621	AA295996	dbEST
Sequence 570	AA182409	dbEST	Sequence 622	AA297432	dbEST
Sequence 571	AA186538	dbEST	Sequence 623	AA298085	dbEST

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Sequence 626	AA303257	dbEST	Sequence 679	AA418681	dbEST
Sequence 627	AA303237 AA304979	dbEST	Sequence 680	AA419263	dbEST
Sequence 628	AA304979 AA305319	dbEST	Sequence 681	AA421682	dbEST
Sequence 629		dbEST	Sequence 682	AA423943	dbEST
Sequence 630	AA305494	dbEST	Sequence 683	AA425218	dbEST
Sequence 631	AA306772	dbEST	•	AA425619	dbEST
Sequence 632	AA307286		Sequence 684	AA428329	dbEST
Sequence 633	AA307539	dbEST	Sequence 685	AA429721	dbEST
Sequence 634	AA307857	dbEST	Sequence 686	AA430381	dbEST
Sequence 635	AA309002	dbEST	Sequence 687	AA430400	dbEST
Sequence 636	AA309127	dbEST	Sequence 688	AA431226	dbEST
Sequence 637	AA310469	dbEST	Sequence 689	AA431220 AA432075	dbEST
Sequence 638	AA311028	dbEST	Sequence 690	AA435887	dbEST
Sequence 639	AA312078	dbEST	Sequence 691		dbEST
Sequence 640	AA312962	dbEST	Sequence 692	AA436765 AA443154	
Sequence 641	AA313164	dbEST	Sequence 693		dbEST dbEST
Sequence 642	AA313653	dbEST	Sequence 694	AA443806	
Sequence 643	AA313692	dbEST	Sequence 695	AA443870	dbEST
Sequence 644	AA313922	dbEST	Sequence 696	AA447116	dbEST
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Sequence 674	AA410580	dbEST	Sequence 726	AA492272	dbEST
Sequence 675	AA411736	dbEST	Sequence 727	AA492280	dbEST
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Sequence 747	AA531563	dbEST	Sequence 799	AA631206	dbEST
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Sequence 749	AA533575	dbEST	Sequence 801		dbEST
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Sequence 760	AA554718	dbEST	Sequence 812	AA687495	dbEST
Sequence 761	AA555227	dbEST	Sequence 813	AA702422	dbEST
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Sequence 779	AA583997	dbEST	Sequence 831	AA774523	dbEST
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Sequence 834	AA779728	dbEST	Sequence 886	AA992596	dbEST
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Sequence 850	AA861631	dbEST	Sequence 902	AI033904	dbEST
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Sequence 987	Al281509	dbEST	Sequence 1039	AI557495	dbEST

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Sequence 1040	Al557599	dbEST	Sequence 1092	A1753269	dbEST
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Sequence 1144	AW027250	dbEST	Sequence 1196	N99698	dbEST
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Sequence 1195		dbEST	Sequence 1245	AB002387	GENBANK
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Sequence 1297	AF020736	GENBANK	Sequence 1349	AF077045	GENBANK

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Sequence 1353 AF078847 GENBANK Sequence 1406 AF195417 GENBANK Sequence 1354 AF078860 GENBANK Sequence 1406 AF201077 GENBANK Sequence 1355 AF078862 GENBANK Sequence 1407 AJ004913 GENBANK Sequence 1356 AF08457 GENBANK Sequence 1407 AJ004913 GENBANK Sequence 1356 AF08457 GENBANK Sequence 1407 AJ004913 GENBANK Sequence 1356 AF085845 GENBANK Sequence 1410 AJ007388 GENBANK Sequence 1359 AF085850 GENBANK Sequence 1410 AJ0070714 GENBANK Sequence 1359 AF085850 GENBANK Sequence 1410 AJ0070714 GENBANK Sequence 1359 AF085850 GENBANK Sequence 1411 AJ010071 GENBANK Sequence 1360 AF086712 GENBANK Sequence 1412 AJ204172 GENBANK Sequence 1361 AF086336 GENBANK Sequence 1413 AJ224172 GENBANK Sequence 1363 AF086351 GENBANK Sequence 1414 AJ224819 GENBANK Sequence 1363 AF086351 GENBANK Sequence 1416 AJ228139 GENBANK Sequence 1364 AF086481 GENBANK Sequence 1416 AJ228139 GENBANK Sequence 1365 AF086567 GENBANK Sequence 1416 AJ228139 GENBANK Sequence 1366 AF08657 GENBANK Sequence 1416 AJ049233 GENBANK Sequence 1366 AF08657 GENBANK Sequence 1416 AJ049233 GENBANK Sequence 1366 AF086567 GENBANK Sequence 1416 AJ049233 GENBANK Sequence 1366 AF086567 GENBANK Sequence 1418 AJ049252 GENBANK Sequence 1368 AF093668 GENBANK Sequence 1418 AJ049252 GENBANK Sequence 1366 AF098687 GENBANK Sequence 1418 AJ049252 GENBANK Sequence 1366 AF098687 GENBANK Sequence 1414 AJ04923 GENBANK Sequence 1370 AF0998687 GENBANK Sequence 1421 AJ049347 GENBANK Sequence 1370 AF0998687 GENBANK Sequence 1422 AJ049447 GENBANK Sequence 1370 AF099868 GENBANK Sequence 1421 AJ049989 GENBANK Sequence 1371 AF100741 GENBANK Sequence 1422 AJ049447 GENBANK Sequence 1371 AF100741 GENBANK Sequence 1425 AJ050089 GENBANK Sequence 1371 AF100741 GENBANK Sequence 1425 AJ050089 GENBANK Sequence 1371 AF100741 GENBANK Sequence 1427 AJ050041 GENBANK Sequence 1371 AF10265 GENBANK Sequence 1427 AJ050041 GENBANK Sequence 1370 AF10784 GENBANK Sequence 1427 AJ050041 GENBANK Sequence 1370 AF10784 GENBANK Sequence 1427 AJ050049 GENBANK Sequence 1370 AF117764 GENBANK Sequence 1427 AJ050049 GENBANK Sequ						
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Sequence 1400 Al 170074 CENDANIA	•			<u>-</u>		
Sequence 1401 AF188745 GENBANK Sequence 1453 D13266 GENBANK	•			•		
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		D40040	OTNIDANIK	Sequence 1506	D89077	GENBANK
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Sequence 14			GENBANK	•	E01888	GENBANK
Sequence 14			GENBANK	Sequence 1508	E01956	GENBANK
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Sequence 14		_	GENBANK	Sequence 1510	E03569	GENBANK
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Sequence 1		D16480	GENBANK	Sequence 1517	J03799	GENBANK
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Sequence 1	469	D21262	GENBANK	Sequence 1521	J05192	GENBANK
Sequence 1	470	D23662	GENBANK	Sequence 1522	K01911	GENBANK
Sequence 1	471	D25542	GENBANK	Sequence 1523	K03002	GENBANK
Sequence 1	472	D28473	GENBANK	Sequence 1524	L02547	GENBANK
Sequence 1	473	D29954	GENBANK	Sequence 1525	L06070	GENBANK
Sequence 1	474	D30655	GENBANK	Sequence 1526	L07033	GENBANK
Sequence 1		D31767	GENBANK	Sequence 1527	L07261	GENBANK
Sequence 1	476	D31883	GENBANK	Sequence 1528	L07515	GENBANK
Sequence 1	477	D31884	GENBANK	Sequence 1529	L08441	GENBANK
Sequence 1		D31891	GENBANK	Sequence 1530	L11284	GENBANK
Sequence 1	1479	D37991	GENBANK	Sequence 1531	L11667	GENBANK
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Sequence 1	1481	D38293	GENBANK	Sequence 1533	L13806	GENBANK
Sequence 1	1482	D38491	GENBANK	Sequence 1534	L14837	GENBANK
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Sequence 1	1489	D50372	GENBANK	Sequence 1541	L34840	GENBANK
Sequence 1	1490	D50525	GENBANK	Sequence 1542	L37080	GENBANK
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Sequence 1		D63997	GENBANK	Sequence 1546	L42110	GENBANK
Sequence 1	1495	D82345	GENBANK	Sequence 1547	L47647	GENBANK
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Sequence 1	1498	D86971	GENBANK	Sequence 1550	M10036	GENBANK
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Sequence 1		D86985	GENBANK	Sequence 1552	M11119	GENBANK
Sequence 1		D87078	GENBANK	Sequence 1553	M13231	GENBANK
Sequence 1		D87666	GENBANK	Sequence 1554	M13692	GENBANK
Sequence 1		D87667	GENBANK	Sequence 1555	M14200	GENBANK
Sequence 1		D88674	GENBANK	Sequence 1556	M15885	GENBANK
Sequence 1		D89053	GENBANK	Sequence 1557	M15887	GENBANK
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Sequence 1		M16804	GENBANK	Sequence 1612	U17714	GENBANK
Sequence 1		M17323	GENBANK	Sequence 1613	U18914	GENBANK
Sequence 1		M17885	GENBANK	Sequence 1614	U21858	GENBANK
Sequence 1		M20030	GENBANK		U24169	GENBANK
Sequence 1		M20372	GENBANK	Sequence 1615	U25789	GENBANK
Sequence 1		M21154	GENBANK	Sequence 1616	U28249	GENBANK
Sequence 1		M21895	GENBANK	Sequence 1617	U30826	GENBANK
Sequence 1		M21896	GENBANK	Sequence 1618	U31905	GENBANK
Sequence 1		M22382	GENBANK	Sequence 1619	U33821	GENBANK
Sequence 1		M22538	GENBANK	Sequence 1620		GENBANK
Sequence 1		M22590	GENBANK	Sequence 1621	U38654	
Sequence 1		M22920	GENBANK	Sequence 1622	U38784	GENBANK GENBANK
Sequence 1		M23613	GENBANK	Sequence 1623	U39360	GENBANK
Sequence 1	1572	M24194	GENBANK	Sequence 1624	U41515	GENBANK
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Sequence 1		M26663	GENBANK	Sequence 1626	U48857	GENBANK GENBANK
Sequence 1		M27335	GENBANK	Sequence 1627	U51004	GENBANK
Sequence 1	1576	M27937	GENBANK	Sequence 1628	U51920	
Sequence 1	1577	M28016	GENBANK	Sequence 1629	U66469	GENBANK
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Sequence 1	1579	M31627	GENBANK	Sequence 1631	U68140	GENBANK
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Sequence 1	1583	M55421	GENBANK	Sequence 1635	U78516	GENBANK
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Sequence '	1585	M59849	GENBANK	Sequence 1637	U80034	GENBANK
Sequence '	1586	M60457	GENBANK	Sequence 1638	U80456	GENBANK
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Sequence 1	1589	M64347	GENBANK	Sequence 1641	U97280	GENBANK
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Sequence	1591	M74777	GENBANK	Sequence 1643	X02544	GENBANK
Sequence	1592	M82882	GENBANK	Sequence 1644	X04707	GENBANK
Sequence	1593	M83822	GENBANK	Sequence 1645	X05332	GENBANK
Sequence	1594	M84443	GENBANK	Sequence 1646	X06747	GENBANK
Sequence	1595	M88163	GENBANK	Sequence 1647	X13238	GENBANK
Sequence		M90104	GENBANK	Sequence 1648	X13839	GENBANK
Sequence		M90309	GENBANK	Sequence 1649	X15183	GENBANK
Sequence		M97856	GENBANK	Sequence 1650	X15187	GENBANK
Sequence		S39329	GENBANK	Sequence 1651	X15722	GENBANK
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Sequence		S70154	GENBANK	Sequence 1654	X52943	GENBANK
Sequence		S73591	GENBANK	Sequence 1655	X53280	GENBANK
Sequence		S78203	GENBANK	Sequence 1656	X53605	GENBANK
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Sequence		S94541	GENBANK	Sequence 1659	X60656	GENBANK
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Sequence		U07919	GENBANK	Sequence 1661	X61100	GENBANK
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Sequence 1664	X74070	GENBANK	Sequence 1717	AA037216	dbEST
Sequence 1665		GENBANK	Sequence 1718	AA037828	dbEST
Sequence 1666	X74331	GENBANK	Sequence 1719	AA037877	dbEST
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Sequence 1668	X76013	GENBANK	Sequence 1721	AA040820	dbEST
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Sequence 1771	AA126373	dbEST	Sequence 1823	AA191661	dbEST
Sequence 1772	AA127336	dbEST	Sequence 1824	AA193247	dbEST
Sequence 1773	AA127369	dbEST	Sequence 1825	AA195617	dbEST
Sequence 1774	AA128114	dbEST	Sequence 1826	AA195831	dbEST
Sequence 1775	AA130162	dbEST	Sequence 1827	AA196978	dbEST
Sequence 1776	AA130229	dbEST	Sequence 1828	AA203492	dbEST
Sequence 1777	AA130262	dbEST	Sequence 1829	AA204854	dbEST
Sequence 1778	AA130784	dbEST	Sequence 1830	AA205647	dbEST
Sequence 1779	AA131198	dbEST	Sequence 1831	AA205870	dbEST
Sequence 1780	AA131749	dbEST	Sequence 1832	AA206137	dbEST
Sequence 1781	AA131961	dbEST	Sequence 1833	AA206214	dbEST
Sequence 1782	AA132508	dbEST	Sequence 1834	AA207266	dbEST
Sequence 1783	AA132633	dbEST	Sequence 1835	AA210741	dbEST
Sequence 1784	AA132992	dbEST	Sequence 1836	AA210833	dbEST
Sequence 1785	AA133692	dbEST	Sequence 1837	AA210840	dbEST
Sequence 1786	AA134266	dbEST	Sequence 1838	AA218536	dbEST
Sequence 1787	AA134267	dbEST	Sequence 1839	AA223774	dbEST
Sequence 1788	AA134527	dbEST	Sequence 1840	AA224132	dbEST
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Sequence 1791	AA136874	dbEST	Sequence 1843	AA225002	dbEST
Sequence 1792	AA143438	dbEST	Sequence 1844	AA225458	dbEST
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Sequence 1795	AA147507	dbEST	Sequence 1847	AA226275	dbEST
Sequence 1796	AA148019	dbEST	Sequence 1848	AA226321	dbEST
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Sequence 1798	AA156782	dbEST	Sequence 1850	AA226709	dbEST
Sequence 1799	AA156862	dbEST	Sequence 1851	AA227873	dbEST
Sequence 1800	AA157741	dbEST	Sequence 1852	AA228440	dbEST
Sequence 1801	AA159774	dbEST	Sequence 1853	AA229199	dbEST
Sequence 1802	AA159849	dbEST	Sequence 1854	AA229424	dbEST
Sequence 1803	AA159910	dbEST	Sequence 1855	AA229611	dbEST
Sequence 1804	AA165027	dbEST	Sequence 1856	AA230169	dbEST
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Sequence 1809	AA167766	dbEST	Sequence 1861	AA234818	dbEST
Sequence 1810	AA167826	dbEST	Sequence 1862	AA235370	dbEST
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Sequence 1812	AA169736	dbEST	Sequence 1864	AA236667	dbEST
Sequence 1813	AA173331	dbEST	Sequence 1865	AA236904	dbEST
Sequence 1814	AA173998	dbEST	Sequence 1866	AA242864	dbEST
Sequence 1815	AA180137	dbEST	Sequence 1867	AA243230	dbEST
Sequence 1816	AA182948	dbEST	Sequence 1868	AA243980	dbEST
Sequence 1817	AA187288	dbEST	Sequence 1869	AA244052	dbEST
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Sequence 1870	AA249154	dbEST	Sequence 1922	AA313170	dbEST
Sequence 1871	AA251321	dbEST	Sequence 1923	AA313534	dbEST
Sequence 1872	AA253384	dbEST	Sequence 1924	AA313779	dbEST
Sequence 1873	AA256591	dbEST	Sequence 1925	AA314350	dbEST
Sequence 1874	AA258357	dbEST	Sequence 1926	AA314599	dbEST
Sequence 1875	AA259062	dbEST	Sequence 1927	AA314673	dbEST
Sequence 1876	AA261873	dbEST	Sequence 1928	AA315049	dbEST
Sequence 1877	AA262345	dbEST	Sequence 1929	AA315189	dbEST
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Sequence 1878	AA280848	dbEST	Sequence 1931	AA316181	dbEST
Sequence 1879	AA283904	dbEST	Sequence 1932	AA316632	dbEST
Sequence 1880	AA284835	dbEST	Sequence 1933	AA317952	dbEST
Sequence 1881	AA284971	dbEST	Sequence 1934	AA318100	dbEST
Sequence 1882	AA285040	dbEST	Sequence 1935	AA320611	dbEST
Sequence 1883	AA285290	dbEST	Sequence 1936	AA322728	dbEST
Sequence 1884	AA290591	dbEST	Sequence 1937	AA322922	dbEST
Sequence 1885	AA291680	dbEST	Sequence 1938	AA324683	dbEST
Sequence 1886	AA291000 AA292131	dbEST	Sequence 1939	AA325108	dbEST
Sequence 1887		dbEST	Sequence 1940	AA325220	dbEST
Sequence 1888	AA292866		Sequence 1941	AA329110	dbEST
Sequence 1889	AA293802	dbEST	Sequence 1942	AA331359	dbEST
Sequence 1890	AA296465	dbEST	Sequence 1943	AA332139	dbEST
Sequence 1891	AA297215	dbEST	Sequence 1944	AA332551	dbEST
Sequence 1892	AA297512	dbEST	Sequence 1945	AA336798	dbEST
Sequence 1893	AA298596	dbEST		AA340632	dbEST
Sequence 1894	AA298786	dbEST	Sequence 1946	AA342231	dbEST
Sequence 1895	AA298794	dbEST	Sequence 1947	AA344376	dbEST
Sequence 1896	AA299594	dbEST	Sequence 1948	AA345904	dbEST
Sequence 1897	AA300728	dbEST	Sequence 1949	AA347340	dbEST
Sequence 1898	AA301391	dbEST	Sequence 1950	AA347340 AA351799	dbEST
Sequence 1899	AA301907	dbEST	Sequence 1951	AA351799 AA354915	dbEST
Sequence 1900	AA303172	dbEST	Sequence 1952	AA354913 AA355119	dbEST
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Sequence 1902	AA303796	dbEST	Sequence 1954	AA357593 AA358169	dbEST
Sequence 1903	AA303880	dbEST	Sequence 1955	AA356109 AA360190	dbEST
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Sequence 1907	AA305331	dbEST	Sequence 1959		dbEST
Sequence 1908	AA305409	dbEST	Sequence 1960	AA366269	
Sequence 1909	AA305526	dbEST	Sequence 1961	AA369521	dbEST
Sequence 1910	AA305566	dbEST	Sequence 1962	AA370110	dbEST
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Sequence 1912	AA306812	dbEST	Sequence 1964	AA370337	dbEST
Sequence 1913	AA307697	dbEST	Sequence 1965	AA373079	dbEST
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Sequence 1915	AA307849	dbEST	Sequence 1967	AA375298	dbEST
Sequence 1916	AA308051	dbEST	Sequence 1968	AA375312	dbEST
Sequence 1917	AA308273	dbEST	Sequence 1969	AA376292	dbEST
Sequence 1918	AA308332	dbEST	Sequence 1970	AA382481	dbEST
Sequence 1919	AA310057	dbEST	Sequence 1971	AA384322	dbEST
Sequence 1920	AA311505	dbEST	Sequence 1972	AA384529	dbEST
Sequence 1921	AA312652	dbEST	Sequence 1973	AA384756	dbEST

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Common 4074	AA385573	dbEST	Sequence 2026	AA451633	dbEST
Sequence 1974	AA386056	dbEST	Sequence 2027	AA453310	dbEST
Sequence 1975	AA386087	dbEST	Sequence 2028	AA453562	dbEST
Sequence 1976	AA393080	dbEST	Sequence 2029	AA453924	dbEST
Sequence 1977	AA393699	dbEST	Sequence 2030	AA455235	dbEST
Sequence 1978	AA393699 AA397460	dbEST	Sequence 2031	AA456344	dbEST
Sequence 1979		dbEST	Sequence 2032	AA458516	dbEST
Sequence 1980	AA398619	dbEST	Sequence 2033	AA459321	dbEST
Sequence 1981	AA400692		Sequence 2034	AA459517	dbEST
Sequence 1982	AA402000	dbEST	Sequence 2035	AA461128	dbEST
Sequence 1983	AA402238	dbEST	Sequence 2036	AA464133	dbEST
Sequence 1984	AA404669	dbEST	Sequence 2037	AA464338	dbEST
Sequence 1985	AA406168	dbEST	Sequence 2038	AA464568	dbEST
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Sequence 1987	AA418628	dbEST	Sequence 2039	AA468839	dbEST
Sequence 1988	AA418685	dbEST	Sequence 2040		dbEST
Sequence 1989	AA418824	dbEST	Sequence 2041	AA469199	dbEST
Sequence 1990	AA419011	dbEST	Sequence 2042	AA469406	dbEST
Sequence 1991	AA420625	dbEST	Sequence 2043	AA476263	dbEST
Sequence 1992	AA420645	dbEST	Sequence 2044	AA477173	dbEST
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Sequence 1994	AA420826	dbEST	Sequence 2046		dbEST
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Sequence 2003	AA428143	dbEST	Sequence 2055	AA488280	dbEST dbEST
Sequence 2004	AA429173	dbEST	Sequence 2056	AA489032	
Sequence 2005	AA429190	dbEST	Sequence 2057	AA489410	dbEST
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Sequence 2011	AA436395	dbEST	Sequence 2063	AA492280	dbEST
Sequence 2012	AA436730	dbEST	Sequence 2064	AA493512	dbEST
Sequence 2013	AA437224	dbEST	Sequence 2065	AA493742	dbEST
Sequence 2014	AA442145	dbEST	Sequence 2066	AA493962	dbEST
Sequence 2015	AA442287	dbEST	Sequence 2067	AA494344	dbEST
Sequence 2016	AA442415	dbEST	Sequence 2068	AA501752	dbEST
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Sequence 2085	AA513940	dbEST	Sequence 2137	AA573975	dbEST
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Sequence 2091	AA522849	dbEST	Sequence 2143	AA578577	dbEST
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Sequence 2101	AA526430	dbEST	Sequence 2153	AA592908	dbEST
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Sequence 2103	AA526869	dbEST	Sequence 2155	AA599669	dbEST
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Sequence 2199	AA682252	dbEST	Sequence 2251	AA838410	dbEST
Sequence 2200	AA682283	dbEST	Sequence 2252	AA838681	dbEST
Sequence 2201	AA687216	dbEST	Sequence 2253	AA843326	dbEST
Sequence 2202	AA687336	dbEST	Sequence 2254	AA843592	dbEST
Sequence 2203	AA700232	dbEST	Sequence 2255	AA843706	dbEST
Sequence 2204	AA700876	dbEST	Sequence 2256	AA844070	dbEST
Sequence 2205	AA702888	dbEST	Sequence 2257	AA847656	dbEST
Sequence 2206	AA703396	dbEST	Sequence 2258	AA848129	dbEST
Sequence 2207	AA705433	dbEST	Sequence 2259	AA853183	dbEST
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Sequence 2209	AA707607	dbEST	Sequence 2261	AA857476	dbEST
Sequence 2210	AA720578	dbEST	Sequence 2262	AA857936	dbEST
Sequence 2211	AA721375	dbEST	Sequence 2263	AA872606	dbEST
Sequence 2212	AA723525	dbEST	Sequence 2264	AA873182	dbEST
Sequence 2213	AA725400	dbEST	Sequence 2265	AA876668	dbEST
Sequence 2214	AA730534	dbEST	Sequence 2266	AA876897	dbEST
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Sequence 2216	AA731556	dbEST	Sequence 2268	AA878212	dbEST
Sequence 2217	AA731700	dbEST	Sequence 2269	AA883580	dbEST
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Sequence 2231	AA769722	dbEST	Sequence 2283	AA926992	dbEST
Sequence 2232	AA774869	dbEST	Sequence 2284	AA932252	dbEST
Sequence 2233	AA775918	dbEST	Sequence 2285	AA934368	dbEST

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Sequence 2422	Al275175	dbEST	Sequence 2474	AI536024	dbEST dbEST
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Sequence 2434	Al318280	dbEST	Sequence 2486	AI557599	dbEST
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Sequence 2441	Al345416	dbEST	Sequence 2493	AI580162	dbEST

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Sequence 2496	AI591088	dbEST	Sequence 2548	AI770098	dbEST
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,	AB002351 AB002365	GENBANK	Sequence 2868	AB023205	GENBANK
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Sequence 3047	AJ012499	GENBANK	Sequence 3099	D29012	GENBANK
Sequence 3048	AJ132694	GENBANK	Sequence 3100	D29643	GENBANK
Sequence 3049	AJ242829	GENBANK	Sequence 3101	D31764	GENBANK
Sequence 3050	AJ243874	GENBANK	Sequence 3102	D31886	GENBANK
Sequence 3051	AL049229	GENBANK	Sequence 3103	D38073	GENBANK
Sequence 3052	AL049784	GENBANK	Sequence 3104	D38551	GENBANK
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Sequence 3054	AL050037	GENBANK	Sequence 3106	D42047	GENBANK
Sequence 3055	AL050101	GENBANK	Sequence 3107	D42063	GENBANK
Sequence 3056	AL050136	GENBANK	Sequence 3108	D43950	GENBANK
Sequence 3057	AL050265	GENBANK	Sequence 3109	D45370	GENBANK
Sequence 3058	AL050274	GENBANK	Sequence 3110	D49387	GENBANK
Sequence 3059	AL050282	GENBANK	Sequence 3111	D50371	GENBANK
Sequence 3060	AL050373	GENBANK	Sequence 3112	D50372	GENBANK
Sequence 3061	AL079279	GENBANK	Sequence 3113	D55696	GENBANK
Sequence 3062	AL080135	GENBANK	Sequence 3114	D63475	GENBANK
Sequence 3063	AL080172	GENBANK	Sequence 3115	D63478	GENBANK
Sequence 3064	AL080192	GENBANK	Sequence 3116	D63997	GENBANK

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Sequence 3117	D67029 D79991	GENBANK	Sequence 3170	L11566	GENBANK
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Sequence 3119	D83032	GENBANK	Sequence 3172	L13434	GENBANK
Sequence 3120	D83703	GENBANK	Sequence 3173	L13852	GENBANK
Sequence 3121	D83703 D84476	GENBANK	Sequence 3174	L15203	GENBANK
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Sequence 3131	D87666	GENBANK	Sequence 3184	L28010	GENBANK
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Sequence 3167	L07033	GENBANK	Sequence 3219	M26663	GENBANK
Sequence 3168	L08044	GENBANK	Sequence 3220	M28016	GENBANK
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Sequence 3262	S63912	GENBANK	Sequence 3314	U63289	GENBANK
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•	U04209	GENBANK	Sequence 3323	U73824	GENBANK
Sequence 3271	U08815	GENBANK	Sequence 3324	U75272	GENBANK
Sequence 3272	000013	GLIADVIAV	Ocquence 0024	UIULIL	JE: 10/ 1/11

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Sequence 3325	U77720	GENBANK	Sequence 3378	X75304	GENBANK
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Sequence 3686	AA292179	dbEST	Sequence 3738	AA308533	dbEST
Sequence 3687	AA292739	dbEST	Sequence 3739	AA308942	dbEST
Sequence 3688	AA292796	dbEST	Sequence 3740	AA308978	dbEST
acdaeure 2000	MAZJZ130	UDEGI	Dequence 3/40	17/10/00/10	UNLUI

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0 0744	A A 200 4 4 F	abECT.	Sequence 3793	AA318544	dbEST
Sequence 3741	AA309445	dbEST dbEST	Sequence 3794	AA319139	dbEST
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Sequence 3744	AA310181	dbEST	Sequence 3797	AA322246	dbEST
Sequence 3745	AA310292	dbEST	Sequence 3798	AA324251	dbEST
Sequence 3746	AA310460	dbEST	Sequence 3799	AA324411	dbEST
Sequence 3747	AA310561		Sequence 3800	AA324705	dbEST
Sequence 3748	AA310591	dbEST dbEST	Sequence 3801	AA324831	dbEST
Sequence 3749	AA310928		Sequence 3802	AA325221	dbEST
Sequence 3750	AA311044	dbEST	Sequence 3803	AA325658	dbEST
Sequence 3751	AA311227	dbEST	Sequence 3804	AA326793	dbEST
Sequence 3752	AA312177	dbEST	Sequence 3805	AA327228	dbEST
Sequence 3753	AA312242	dbEST	· · · · · · · · · · · · · · · · · · ·	AA327546	dbEST
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Sequence 3759	AA313418	dbEST	Sequence 3811	AA332667	dbEST
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Sequence 3762	AA313837	dbEST	Sequence 3814	AA333526	dbEST
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Sequence 3766	AA314181	dbEST	Sequence 3818	AA336740	dbEST
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Sequence 3790	AA317096	dbEST	Sequence 3842	AA359514	dbEST
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Sequence 3792	AA317956	dbEST	Sequence 3844	AA362778	dbEST

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Sequence 3896	AA424658	dbEST	Gequence 3940	, , , , , , , , , , , , , , , , , , , ,	

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Sequence 3988	AA506299	dbEST	Sequence 4040	AA570171	dbEST
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Sequence 4208	Al039227	dbEST	Sequence 4260	AI167801	dbEST

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Sequence 4311	Al267596	dbEST	Sequence 4363	Al394498	dbEST
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Sequence 4368	Al434639	dbEST	Sequence 4420	Al690236	dbEST
Sequence 4369	Al436714	dbEST	Sequence 4421	Al692940	dbEST
Sequence 4370	Al439173	dbEST	Sequence 4422	Al692995	dbEST
Sequence 4371	Al453358	dbEST	Sequence 4423	Al694572	dbEST
Sequence 4372	Al473779	dbEST	Sequence 4424	Al697815	dbEST
Sequence 4373	Al476254	dbEST	Sequence 4425	Al701119	dbEST
Sequence 4374	Al499378	dbEST	Sequence 4426	AI708995	dbEST
Sequence 4375	Al499439	dbEST	Sequence 4427	Al732541	dbEST
Sequence 4376	Al499933	dbEST	Sequence 4428	Al732557	dbEST
Sequence 4377	Al521270	dbEST	Sequence 4429	Al744451	dbEST
Sequence 4378	AI525426	dbEST	Sequence 4430	AI745280	dbEST
Sequence 4379	AI525773	dbEST	Sequence 4431	AI751583	dbEST
Sequence 4380	AI525843	dbEST	Sequence 4432	Al751914	dbEST
Sequence 4381	Al537498	dbEST	Sequence 4433	AI752822	dbEST
Sequence 4382	AI538681	dbEST	Sequence 4434	AI753269	dbEST
Sequence 4383	AI547284	dbEST	Sequence 4435	AI754013	dbEST
Sequence 4384	Al547309	dbEST	Sequence 4436	AI755085	dbEST
Sequence 4385	Al557226	dbEST	Sequence 4437	Al761508	dbEST
Sequence 4386	Al557495	dbEST	Sequence 4438	Al761728	dbEST
Sequence 4387	AI557626	dbEST	Sequence 4439	AI768848	dbEST
Sequence 4388	AI559999	dbEST	Sequence 4440	AI769145	dbEST
Sequence 4389	Al566084	dbEST	Sequence 4441	AI791719	dbEST
Sequence 4390	Al566490	dbEST	Sequence 4442	AI802998	dbEST
Sequence 4391	Al570942	dbEST	Sequence 4443	AI806161	dbEST
Sequence 4392	AI571069	dbEST	Sequence 4444	AI806996	dbEST
Sequence 4393	AI580160	dbEST	Sequence 4445	AI815416	dbEST
Sequence 4394	AI587118	dbEST	Sequence 4446	AI815449	dbEST
Sequence 4395	AI588087	dbEST	Sequence 4447	AI815497	dbEST
Sequence 4396	AI589633	dbEST	Sequence 4448	Al815829	dbEST
Sequence 4397	Al608968	dbEST	Sequence 4449	AI815966	dbEST
Sequence 4398	Al628609	dbEST	Sequence 4450	AI815976	dbEST
Sequence 4399	Al630282	dbEST	Sequence 4451	AI816033	dbEST
Sequence 4400	Al636577	dbEST	Sequence 4452	AI820995	dbEST
Sequence 4401	Al636911	dbEST	Sequence 4453	Al829303	dbEST
Sequence 4402	Al638327	dbEST	Sequence 4454	Al858066	dbEST
Sequence 4403	Al648682	dbEST	Sequence 4455	Al860714	dbEST
Sequence 4404	Al651069	dbEST	Sequence 4456	Al866262	dbEST
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Sequence 4406	Al651976	dbEST	Sequence 4458	Al872085	dbEST
Sequence 4407	Al652084	dbEST	Sequence 4459	Al878878	dbEST
Sequence 4408	AI652555	dbEST	Sequence 4460	AI879010	dbEST
Sequence 4409	Al660243	dbEST	Sequence 4461	AI879059	dbEST
Sequence 4410	Al669253	dbEST	Sequence 4462	AI879242	dbEST
Sequence 4411	Al670002	dbEST	Sequence 4463	Al884578	dbEST
Sequence 4412	AI672362	dbEST	Sequence 4464	Al885833	dbEST
Sequence 4413	Al674583	dbEST	Sequence 4465	Al886418	dbEST
Sequence 4414	Al675889	dbEST	Sequence 4466	Al887517	dbEST
Sequence 4415	Al679629	dbEST	Sequence 4467	Al887664	dbEST
Sequence 4416	Al683183	dbEST	Sequence 4468	Al911703	dbEST

Sequence 4469	Al914133	dbEST	Sequence 4521	AW026680	dbEST
Sequence 4470	Al917649	dbEST	Sequence 4522	AW027007	dbEST
Sequence 4471	Al924598	dbEST	Sequence 4523	AW027219	dbEST
Sequence 4472	Al929113	dbEST	Sequence 4524	AW029146	dbEST
Sequence 4473	Al929453	dbEST	Sequence 4525	AW058559	dbEST
Sequence 4474	Al934153	dbEST	Sequence 4526	AW062376	dbEST
Sequence 4475	Al934595	dbEST	Sequence 4527	AW071906	dbEST
Sequence 4476	Al935291	dbEST	Sequence 4528	AW075433	dbEST
Sequence 4477	Al935816	dbEST	Sequence 4529	AW084950	dbEST
Sequence 4478	Al937060	dbEST	Sequence 4530	AW129510	dbEST
Sequence 4479	Al951021	dbEST	Sequence 4531	AW132009	dbEST
Sequence 4480	AI951536	dbEST	Sequence 4532	AW134746	dbEST
Sequence 4481	Al954800	dbEST	Sequence 4533	AW134867	dbEST
Sequence 4482	AI962736	dbEST	Sequence 4534	AW151254	dbEST
Sequence 4483	Al973126	dbEST	Sequence 4535	AW152107	dbEST
Sequence 4484	Al973218	dbEST	Sequence 4536	AW160669	dbEST
Sequence 4485	Al983430	dbEST	Sequence 4537	AW161467	dbEST
Sequence 4486	Al991281	dbEST	Sequence 4538	AW161705	dbEST
Sequence 4487	AL035886	dbEST	Sequence 4539	AW163208	dbEST
Sequence 4488	AL035985	dbEST	Sequence 4540	AW163441	dbEST
Sequence 4489	AL036026	dbEST	Sequence 4541	AW167481	dbEST
Sequence 4490	AL036415	dbEST	Sequence 4542	AW169941	dbEST
Sequence 4491	AL036445	dbEST	Sequence 4543	AW170485	dbEST
Sequence 4492	AL036763	dbEST	Sequence 4544	AW176062	dbEST
Sequence 4493	AL036764	dbEST	Sequence 4545	AW177936	dbEST
Sequence 4494	AL037724	dbEST	Sequence 4546	AW177948	dbEST
Sequence 4495	AL038058	dbEST	Sequence 4547	AW177957	dbEST
Sequence 4496	AL038660	dbEST	Sequence 4548	AW179006	dbEST
Sequence 4497	AL038721	dbEST	Sequence 4549	AW181932	dbEST
Sequence 4498	AL039517	dbEST	Sequence 4550	AW183601	dbEST
Sequence 4499	AL039691	dbEST	Sequence 4551	AW188666	dbEST
Sequence 4500	AL040505	dbEST	Sequence 4552	C02912	dbEST
Sequence 4501	AL040993	dbEST	Sequence 4553	C16620	dbEST
Sequence 4502	AL041470	dbEST	Sequence 4554	C18148	dbEST
Sequence 4503	AL042896	dbEST	Sequence 4555	C18436	dbEST
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Sequence 4509	AL046313	dbEST	Sequence 4561	F19390	dbEST
Sequence 4510	AL047277	dbEST	Sequence 4562	F25339	dbEST
Sequence 4511	AL047334	dbEST	Sequence 4563	F27302	dbEST
Sequence 4512	AL047838	dbEST	Sequence 4564	F27796	dbEST
Sequence 4513	AL048554	dbEST	Sequence 4565	F28190	dbEST
Sequence 4514	AL110463	dbEST	Sequence 4566	F34889	dbEST
Sequence 4515	AL118956	dbEST	Sequence 4567	H00112	dbEST
Sequence 4516	AL120807	dbEST	Sequence 4568	H01419	dbEST
Sequence 4517	AW006392	dbEST	Sequence 4569	H01979	dbEST
Sequence 4518	AW014590	dbEST	Sequence 4570	H02725	dbEST
Sequence 4519	AW016611	dbEST	Sequence 4571	H06040	dbEST
Sequence 4520	AW026543	dbEST	Sequence 4572	H06313	dbEST

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Sequence 4573	H08929	dbEST	Sequence 4625	R70116	dbEST
Sequence 4574	H10932	dbEST	Sequence 4626	R77854	dbEST
Sequence 4575	H12090	dbEST	Sequence 4627	R85291	dbEST
Sequence 4576	H12254	dbEST	Sequence 4628	T30604	dbEST
Sequence 4577	H17195	dbEST	Sequence 4629	T59793	dbEST
Sequence 4578	H19153	dbEST	Sequence 4630	T62753	dbEST
Sequence 4579	H19367	dbEST	Sequence 4631	T65174	dbEST
Sequence 4580	H20888	dbEST	Sequence 4632	T65642	dbEST
Sequence 4581	H23370	dbEST	Sequence 4633	T68057	dbEST
Sequence 4582	H25211	dbEST	Sequence 4634	T71122	dbEST
Sequence 4583	H25825	dbEST	Sequence 4635	T74004	dbEST
Sequence 4584	H26319	dbEST	Sequence 4636	T79957	dbEST
Sequence 4585	H28581	dbEST	Sequence 4637	T81666	dbEST
Sequence 4586	H29149	dbEST	Sequence 4638	T87572	dbEST
Sequence 4587	H29635	dbEST	Sequence 4639	W01739	dbEST
Sequence 4588	H39809	dbEST	Sequence 4640	W02575	dbEST
Sequence 4589	H49903	dbEST	Sequence 4641	W04588	dbEST
Sequence 4590	H54240	dbEST	Sequence 4642	W28466	dbEST
Sequence 4591	H60247	dbEST	Sequence 4643	W31167	dbEST
Sequence 4592	H61232	dbEST	Sequence 4644	W32156	dbEST
Sequence 4593	H70543	dbEST	Sequence 4645	W38407	dbEST
Sequence 4594	H71076	dbEST	Sequence 4646	W38913	dbEST
Sequence 4595	H79973	dbEST	Sequence 4647	W39053	dbEST
Sequence 4596	H83317	dbEST	Sequence 4648	W48827	dbEST
Sequence 4597	H83328	dbEST	Sequence 4649	W72729	dbEST
Sequence 4598	H85200	dbEST	Sequence 4650	W77826	dbEST
Sequence 4599	H87807	dbEST	Sequence 4651	W79340	dbEST
Sequence 4600	H89883	dbEST	Sequence 4652	W80673	dbEST
Sequence 4601	H97090	dbEST	Sequence 4653	W85764	dbEST
Sequence 4602	H99586	dbEST	Sequence 4654	Z33579	dbEST
Sequence 4603	N21201	dbEST	Sequence 4655	N91259	NUCPATENT
Sequence 4604	N24459	dbEST	Sequence 4656	N91825	NUCPATENT
Sequence 4605	N26421	dbEST	Sequence 4657	Q37811	NUCPATENT
Sequence 4606	N28008	dbEST	Sequence 4658	Q65520	NUCPATENT
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Sequence 4608	N32737	dbEST	Sequence 4660	T02792	NUCPATENT
Sequence 4609	N41418	dbEST	Sequence 4661	T60740	NUCPATENT
Sequence 4610	N42929	dbEST	Sequence 4662	T72175	NUCPATENT
Sequence 4611	N47360	dbEST	Sequence 4663	T79274	NUCPATENT
Sequence 4612	N71080	dbEST	Sequence 4664	T97897	NUCPATENT
Sequence 4613	R11885	dbEST	Sequence 4665	V20443	NUCPATENT
Sequence 4614	R13878	dbEST	Sequence 4666	V29267	NUCPATENT
Sequence 4615	R17110	dbEST	Sequence 4667	V35457	NUCPATENT
Sequence 4616	R18274	dbEST	Sequence 4668	V43612	NUCPATENT
Sequence 4617	R24400	dbEST	Sequence 4669	V57903	NUCPATENT
Sequence 4618	R24834	dbEST	Sequence 4670	V58544	NUCPATENT
Sequence 4619	R34553	dbEST	Sequence 4671	V58619	NUCPATENT
Sequence 4620	R36477	dbEST	Sequence 4672	V59096	NUCPATENT
Sequence 4621	R42994	dbEST	Sequence 4673	V59539	NUCPATENT
Sequence 4622	R54273	dbEST	Sequence 4674	V62427	NUCPATENT
Sequence 4623	R59031	dbEST	Sequence 4675	V64158	NUCPATENT
Sequence 4624	R61601	dbEST	Sequence 4676	V69855	NUCPATENT

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Sequence 4677	V70354	NUCPATENT		quence 4726	AB002346	GENBANK
Sequence 4678	V72211	NUCPATENT		quence 4727	AB002353	GENBANK
Sequence 4679	V81414	NUCPATENT		quence 4728	AB002375	GENBANK
Sequence 4680	V84510	NUCPATENT		quence 4729	AB002380	GENBANK
Sequence 4681	V86232	NUCPATENT		quence 4730	AB002386	GENBANK
Sequence 4682	V89565	NUCPATENT		quence 4731	AB007929	GENBANK
Sequence 4683	V90009	NUCPATENT	Se	quence 4732	AB007932	GENBANK
Sequence 4684	X00630	NUCPATENT	Se	quence 4733	AB007956	GENBANK
Sequence 4685	X03841	NUCPATENT	Se	quence 4734	AB011120	GENBANK
Sequence 4686	X07429	NUCPATENT	Se	quence 4735	AB011147	GENBANK
Sequence 4687	X20426	NUCPATENT	Se	quence 4736	AB014888	GENBANK
Sequence 4688	X20437	NUCPATENT	Se	quence 4737	AB016533	GENBANK
Sequence 4689	X27262	NUCPATENT	Se	quence 4738	AB017430	GENBANK
Sequence 4690	X37385	NUCPATENT	Se	quence 4739	AB017971	GENBANK
Sequence 4691	X37553	NUCPATENT	Se	quence 4740	AB018284	GENBANK
Sequence 4692	X39825	NUCPATENT	Se	quence 4741	AB018313	GENBANK
Sequence 4693	X39912	NUCPATENT	Se	quence 4742	AB019392	GENBANK
Sequence 4694	X52268	NUCPATENT	Se	quence 4743	AB019568	GENBANK
Sequence 4695	X56415	NUCPATENT	Se	quence 4744	AB020665	GENBANK
Sequence 4696	X59352	NUCPATENT	Se	quence 4745	AB020679	GENBANK
Sequence 4697	X80676	NUCPATENT	Se	quence 4746	AB023145	GENBANK
Sequence 4698	X87397	NUCPATENT	Se	quence 4747	AB023177	GENBANK
Sequence 4699	X97772	NUCPATENT		quence 4748	AB023201	GENBANK
Sequence 4700	X97790	NUCPATENT	Se	quence 4749	AB027466	GENBANK
Sequence 4701	Z06250	NUCPATENT	Se	quence 4750	AB028624	GENBANK
Sequence 4702	Z15124	NUCPATENT		quence 4751	AB028945	GENBANK
Sequence 4703	Z15132	NUCPATENT	Se	quence 4752	AB029290	GENBANK
Sequence 4704	Z15187	NUCPATENT		quence 4753	AB032951	 GENBANK
Sequence 4705	Z15987	NUCPATENT	Se	quence 4754	AB033033	GENBANK
Sequence 4706	Z24892	NUCPATENT	Se	quence 4755	AB033076	GENBANK
Sequence 4707	Z24893	NUCPATENT	Se	quence 4756	AB033080	GENBANK
Sequence 4708	Z27237	NUCPATENT		quence 4757	AD001528	GENBANK
Sequence 4709	Z33949	NUCPATENT		quence 4758	AF001893	GENBANK
Sequence 4710	AC29436	PREPATNUC		quence 4759	AF002210	GENBANK
Sequence 4711	AC31443	PREPATNUC		quence 4760	AF004813	GENBANK
Sequence 4712	AC31770	PREPATNUC	Se	quence 4761	AF006043	GENBANK
Sequence 4713	AC33242	PREPATNUC		quence 4762	AF007217	GENBANK
Sequence 4714	AC33576	PREPATNUC		quence 4763	AF009615	GENBANK
Sequence 4715	AC33673	PREPATNUC		quence 4764	AF010233	GENBANK
Sequence 4716	AC33992	PREPATNUC	Se	quence 4765	AF010315	GENBANK
Sequence 4717	AC34287	PREPATNUC	Se	quence 4766	AF012872	GENBANK
Sequence 4718	AC34371	PREPATNUC		quence 4767	AF013160	GENBANK
Sequence 4719	AC34444	PREPATNUC		quence 4768	AF015283	GENBANK
Sequence 4720	AC34517	PREPATNUC	Sec	quence 4769	AF017256	GENBANK
Sequence 4721	AC34738	PREPATNUC		quence 4770	AF020202	GENBANK
Sequence 4722	AC34865	PREPATNUC		quence 4771	AF021351	GENBANK
Sequence 4723	AC35363	PREPATNUC		quence 4772	AF028832	GENBANK
0				quence 4773	AF035295	GENBANK
Sequence #	Accession	Database		quence 4774	AF035812	GENBANK
Coguence 4704	#	OFNDANIC		quence 4775	AF038404	GENBANK
Sequence 4724	A06846	GENBANK		quence 4776	AF041483	GENBANK
Sequence 4725	A17546	GENBANK	Sed	quence 4777	AF045167	GENBANK

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Sequence 4778	AF047599	GENBANK	Sequence 483	0 AJ243310	GENBANK
Sequence 4779	AF052187	GENBANK	Sequence 483		GENBANK
Sequence 4780	AF054990	GENBANK	Sequence 483		
Sequence 4781	AF060515	GENBANK	Sequence 483		GENBANK
Sequence 4782	AF061258	GENBANK	Sequence 483		GENBANK
Sequence 4783	AF064769	GENBANK	Sequence 483		GENBANK
Sequence 4784	AF070523	GENBANK	•		GENBANK
Sequence 4785	AF070561	GENBANK	Sequence 483		GENBANK
Sequence 4786	AF070501 AF070651		Sequence 483		GENBANK
•		GENBANK	Sequence 483		GENBANK
Sequence 4787	AF070657	GENBANK	Sequence 483		GENBANK
Sequence 4788	AF070665	GENBANK	Sequence 484		GENBANK
Sequence 4789	AF071202	GENBANK	Sequence 484		GENBANK
Sequence 4790	AF074016	GENBANK	Sequence 484		GENBANK
Sequence 4791	AF077030	GENBANK	Sequence 484		GENBANK
Sequence 4792	AF078858	GENBANK	Sequence 484		GENBANK
Sequence 4793	AF083107	GENBANK	Sequence 484		GENBANK
Sequence 4794	AF084555	GENBANK	Sequence 484		GENBANK
Sequence 4795	AF085243	GENBANK	Sequence 484		GENBANK
Sequence 4796	AF085858	GENBANK	Sequence 484		GENBANK
Sequence 4797	AF089747	GENBANK	Sequence 484		GENBANK
Sequence 4798	AF091083	GENBANK	Sequence 485		GENBANK
Sequence 4799	AF092038	GENBANK	Sequence 485		GENBANK
Sequence 4800	AF104252	GENBANK	Sequence 485	2 D87292	GENBANK
Sequence 4801	AF114816	GENBANK	Sequence 485	3 D87438	GENBANK
Sequence 4802	AF124440	GENBANK	Sequence 485	4 D87666	GENBANK
Sequence 4803	AF131775	GENBANK	Sequence 4859	5 D89675	GENBANK
Sequence 4804	AF131802	GENBANK	Sequence 4856	5 D90373	GENBANK
Sequence 4805	AF132956	GENBANK	Sequence 485	7 D90427	GENBANK
Sequence 4806	AF136450	GENBANK	Sequence 4858	B D90452	GENBANK
Sequence 4807	AF145316	GENBANK	Sequence 4859	9 E02628	GENBANK
Sequence 4808	AF147331	GENBANK	Sequence 4860	E06721	GENBANK
Sequence 4809	AF151840	GENBANK	Sequence 486°	1 J02642	GENBANK
Sequence 4810	AF151844	GENBANK	Sequence 4862	2 J03802	GENBANK
Sequence 4811	AF151870	GENBANK	Sequence 4863	3 J04183	GENBANK
Sequence 4812	AF151902	GENBANK	Sequence 4864	J04759	GENBANK
Sequence 4813	AF151906	GENBANK	Sequence 4868	K01911	GENBANK
Sequence 4814	AF157028	GENBANK	Sequence 4866		GENBANK
Sequence 4815	AF159056	GENBANK	Sequence 4867		GENBANK
Sequence 4816	AF159295	GENBANK	Sequence 4868		GENBANK
Sequence 4817	AF168956	GENBANK	Sequence 4869		GENBANK
Sequence 4818	AF176555	GENBANK	Sequence 4870		GENBANK
Sequence 4819	AF176642	GENBANK	Sequence 487		GENBANK
Sequence 4820	AF182289	GENBANK	Sequence 4872		GENBANK
Sequence 4821	AF188745	GENBANK	Sequence 4873		GENBANK
Sequence 4822	AF188746	GENBANK	Sequence 4874		GENBANK
Sequence 4823	AF191020	GENBANK	Sequence 4875		GENBANK
Sequence 4824	AF191339	GENBANK	Sequence 4876		GENBANK
Sequence 4825	AJ005259	GENBANK	Sequence 4877		GENBANK
Sequence 4826	AJ006470	GENBANK	Sequence 4878		
Sequence 4827	AJ010482	GENBANK	Sequence 4879		GENBANK
Sequence 4828	AJ132583	GENBANK	Sequence 4880		GENBANK
Sequence 4829	AJ224172	GENBANK	Sequence 4881		GENBANK
400100 4050	, WELT 11 L	CLIRDVIAI	Sequence 488 i	M10906	GENBANK

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Sequence 4882	M11560	GENBANK	Sequence 4934	U67171	GENBANK
Sequence 4883	M12125	GENBANK	Sequence 4935	U70063	GENBANK
Sequence 4884	M12623	GENBANK	Sequence 4936	U79254	GENBANK
Sequence 4885	M13692	GENBANK	Sequence 4937	U79457	GENBANK
Sequence 4886	M15395	GENBANK	Sequence 4938	U82226	GENBANK
Sequence 4887	M15885	GENBANK	Sequence 4939	U82256	GENBANK
Sequence 4888	M16247	GENBANK	Sequence 4940	U94586	GENBANK
Sequence 4889	M17885	GENBANK	Sequence 4941	X01037	GENBANK
Sequence 4890	M17886	GENBANK	Sequence 4942	X01630	GENBANK
Sequence 4891	M21895	GENBANK	Sequence 4943	X04098	GENBANK
Sequence 4892	M22538	GENBANK	Sequence 4944	X04106	GENBANK
Sequence 4893	M22918	GENBANK	Sequence 4945	X04408	GENBANK
Sequence 4894	M26325	GENBANK	Sequence 4946	X04526	GENBANK
Sequence 4895	M26663	GENBANK	Sequence 4947	X13839	GENBANK
Sequence 4896	M26880	GENBANK	Sequence 4948	X15729	GENBANK
Sequence 4897	M29064	GENBANK	Sequence 4949	X57352	GENBANK
Sequence 4898	M33308	GENBANK	Sequence 4950	X58141	GENBANK
Sequence 4899	M34840	GENBANK	Sequence 4951	X60036	GENBANK
Sequence 4900	M36647	GENBANK	Sequence 4952	X63380	GENBANK
Sequence 4901	M60255	GENBANK	Sequence 4953	X63422	GENBANK
Sequence 4902	M61832	GENBANK	Sequence 4954	X63564	GENBANK
Sequence 4903	M64098	GENBANK	Sequence 4955	X68277	GENBANK
Sequence 4904	M64241	GENBANK	Sequence 4956	X69711	GENBANK
Sequence 4905	M64571	GENBANK	Sequence 4957	X73874	GENBANK
Sequence 4906	M68840	GENBANK	Sequence 4958	X76302	GENBANK
Sequence 4907	M81757	GENBANK	Sequence 4959	X76732	GENBANK
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Sequence 4909	S50015	GENBANK	Sequence 4961	X84908	GENBANK
Sequence 4910	S73591	GENBANK	Sequence 4962	Y00815	GENBANK
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Sequence 4912	S75755	GENBANK	Sequence 4964	AA010986	dbEST
Sequence 4913	U02570	GENBANK	Sequence 4965	AA011203	dbEST
Sequence 4914	U02619	GENBANK	Sequence 4966	AA026630	dbEST
Sequence 4915	U07643	GENBANK	Sequence 4967	AA029143	dbEST
Sequence 4916	U10248	GENBANK	Sequence 4968	AA033869	dbEST
Sequence 4917	U10439	GENBANK	Sequence 4969	AA034436	dbEST
Sequence 4918	U14968	GENBANK	Sequence 4970	AA037190	dbEST
Sequence 4919	U14970	GENBANK	Sequence 4971	AA040127	dbEST
Sequence 4920	U15174	GENBANK	Sequence 4972	AA043951	dbEST
Sequence 4921	U16738	GENBANK	Sequence 4973	AA044659	dbEST
Sequence 4922	U18197	GENBANK	Sequence 4974	AA056603	dbEST
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Sequence 4925	U25766	GENBANK	Sequence 4977	AA064886	dbEST
Sequence 4926	U34038	GENBANK	Sequence 4978	AA071511	dbEST
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Sequence 4932	U52962	GENBANK	Sequence 4984	AA127185	dbEST
Sequence 4933	U52969	GENBANK	Sequence 4985	AA132342	dbEST
-	-		254451105 4000	. 4 (102072	UDEO 1

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Sequence 4986	AA133798	dbEST	Sequence 5038	AA340719	dbEST
Sequence 4987	AA150891	dbEST	Sequence 5039	AA342969	dbEST
Sequence 4988	AA172239	dbEST	Sequence 5040	AA344067	dbEST
Sequence 4989	AA173348	dbEST	Sequence 5041	AA346600	dbEST
Sequence 4990	AA173948	dbEST	Sequence 5042	AA354527	dbEST
Sequence 4991	AA178907	dbEST	Sequence 5043	AA359763	dbEST
Sequence 4992	AA179409	dbEST	Sequence 5044	AA364498	dbEST
Sequence 4993	AA190890	dbEST	Sequence 5045	AA364667	dbEST
Sequence 4994	AA191092	dbEST	Sequence 5046	AA366518	dbEST
Sequence 4995	AA191476	dbEST	Sequence 5047	AA371017	dbEST
Sequence 4996	AA195734	dbEST	Sequence 5048	AA371205	dbEST
Sequence 4997	AA224751	dbEST	Sequence 5049	AA377218	dbEST
Sequence 4998	AA225488	dbEST	Sequence 5050	AA384125	dbEST
Sequence 4999	AA225517	dbEST	Sequence 5051	AA400213	dbEST
Sequence 5000	AA226735	dbEST	Sequence 5052	AA426235	dbEST
Sequence 5001	AA228953	dbEST	Sequence 5053	AA430597	dbEST
Sequence 5002	AA229225	dbEST	Sequence 5054	AA442615	dbEST
Sequence 5003	AA251182	dbEST	Sequence 5055	AA443184	dbEST
Sequence 5004	AA256401	dbEST	Sequence 5056	AA443463	dbEST
Sequence 5005	AA258916	dbEST	Sequence 5057	AA446877	dbEST
Sequence 5006	AA259025	dbEST	Sequence 5058	AA447700	dbEST
Sequence 5007	AA279054	dbEST	Sequence 5059	AA448617	dbEST
Sequence 5008	AA292866	dbEST	Sequence 5060	AA451733	dbEST
Sequence 5009	AA293885	dbEST	Sequence 5061	AA451988	dbEST
Sequence 5010	AA296534	dbEST	Sequence 5062	AA454962	dbEST
Sequence 5011	AA296690	dbEST	Sequence 5063	AA470833	dbEST
Sequence 5012	AA296846	dbEST	Sequence 5064	AA476509	dbEST
Sequence 5013	AA297270	dbEST	Sequence 5065	AA483887	dbEST
Sequence 5014	AA298593	dbEST	Sequence 5066	AA484751	dbEST
Sequence 5015	AA299209	dbEST	Sequence 5067	AA485767	dbEST
Sequence 5016	AA300065	dbEST	Sequence 5068	AA486551	dbEST
Sequence 5017	AA301096	dbEST	Sequence 5069	AA487556	dbEST
Sequence 5018	AA301666	dbEST	Sequence 5070	AA488567	dbEST
Sequence 5019	AA302919	¹ dbEST	Sequence 5071	AA489498	dbEST
Sequence 5020	AA303594	dbEST	Sequence 5072	AA490893	dbEST
Sequence 5021	AA305409	dbEST	Sequence 5073	AA503943	dbEST
Sequence 5022	AA306378	dbEST	Sequence 5074	AA507306	dbEST
Sequence 5023	AA310441	dbEST	Sequence 5075	AA507878	dbEST
Sequence 5024	AA310561	dbEST	Sequence 5076	AA508861	dbEST
Sequence 5025	AA312444	dbEST	Sequence 5077	AA523982	dbEST
Sequence 5026	AA313200	dbEST	Sequence 5078	AA528123	dbEST
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Sequence 5029	AA314757	dbEST	Sequence 5081	AA535185	dbEST
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Sequence 5033	AA324251	dbEST	Sequence 5085	AA554735	dbEST
Sequence 5034	AA325214	dbEST	Sequence 5086	AA570748	dbEST
Sequence 5035	AA327546	dbEST	Sequence 5087	AA573559	dbEST
Sequence 5036	AA333526	dbEST	Sequence 5088	AA573811	dbEST
Sequence 5037	AA339065	dbEST	Sequence 5089	AA578265	dbEST

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Sequence 5090	AA579412	dbEST	Sequence 5142	Al267502	dbEST
Sequence 5091	AA579998	dbEST	Sequence 5143	Al268362	dbEST
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Sequence 5096	AA633399	dbEST	Sequence 5148	Al288388	dbEST
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Sequence 5098	AA669823	dbEST	Sequence 5150	Al307634	dbEST
Sequence 5099	AA703004	dbEST	Sequence 5151	Al310235	dbEST
Sequence 5100	AA740574	dbEST	Sequence 5152	Al360451	dbEST
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Sequence 5106	AA862855	dbEST	Sequence 5158	Al473779	dbEST
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Sequence 5124	Al128091	dbEST	Sequence 5176	AI749886	dbEST
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Sequence 5131	Al189345	dbEST	Sequence 5183	Al929664	dbEST
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Sequence 5140	Al267216	dbEST	Sequence 5192	AW071906	dbEST
Sequence 5141	Al267282	dbEST	Sequence 5193	AW130325	dbEST

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Sequence 5195 AW177929 dbEST Sequence 5247 Z33671 NUCPATENT Sequence 5196 AW177931 dbEST Sequence 5197 C05816 dbEST Sequence 5198 C18484 dbEST Sequence 5199 D58694 dbEST Sequence 5248 A06846 GENBANK Sequence 5200 F24428 dbEST Sequence 5248 A06919 GENBANK Sequence 5200 F24428 dbEST Sequence 5240 A06919 GENBANK Sequence 5201 F27796 dbEST Sequence 5250 A06925 GENBANK Sequence 5202 H00112 dbEST Sequence 5251 A21185 GENBANK Sequence 5202 H00112 dbEST Sequence 5252 A23013 GENBANK Sequence 5203 H08924 dbEST Sequence 5252 A23013 GENBANK Sequence 5204 H18732 dbEST Sequence 5252 A23013 GENBANK Sequence 5206 H58534 dbEST Sequence 5252 A2300220 GENBANK Sequence 5206 H58534 dbEST Sequence 5253 A8000250 GENBANK Sequence 5206 M85357 dbEST Sequence 5254 A8000584 GENBANK Sequence 5206 M85357 dbEST Sequence 5256 A8002312 GENBANK Sequence 5207 M0017 dbEST Sequence 5256 A8002382 GENBANK Sequence 5208 M86292 dbEST Sequence 5257 A8002382 GENBANK Sequence 5210 R70270 dbEST Sequence 5258 A800286 GENBANK Sequence 5210 R70270 dbEST Sequence 5259 A8004851 GENBANK Sequence 5211 W00471 dbEST Sequence 5259 A8004851 GENBANK Sequence 5211 W00471 dbEST Sequence 5261 A8006834 GENBANK Sequence 5211 W07476 dbEST Sequence 5261 A8006834 GENBANK Sequence 5211 W07476 dbEST Sequence 5261 A8006834 GENBANK Sequence 5211 W37945 dbEST Sequence 5264 A8007892 GENBANK Sequence 5215 W73588 dbEST Sequence 5264 A8007802 GENBANK Sequence 5216 W73588 dbEST Sequence 5264 A8007802 GENBANK Sequence 5216 W73758 dbEST Sequence 5264 A8007802 GENBANK Sequence 5216 W73588 dbEST Sequence 5264 A8007802 GENBANK Sequence 5210 W0747ENT Sequence 5268 A8007802 GENBANK Sequence 5210 W0747ENT Sequence 5267 A8007801 GENBANK GENBANK G	Sequence 5194	AW152107	dbEST	Ş	Sequence 5246	Z17710	NUCPATENT
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Sequence 5295	AB033010	GENBANK	Sequence 5347	AF062318	GENBANK
Sequence 5296	AB033017	GENBANK	Sequence 5348	AF064491	GENBANK
Sequence 5297	AB033070	GENBANK	Sequence 5349	AF064819	GENBANK
Sequence 5298	AB033074	GENBANK	Sequence 5350	AF065388	GENBANK
Sequence 5299	AF007791	GENBANK	Sequence 5351	AF068754	GENBANK
Sequence 5300	AF008443	GENBANK	Sequence 5352	AF069073	GENBANK
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Sequence 5302	AF015283	GENBANK	Sequence 5354	AF070523	GENBANK
Sequence 5303	AF017305	GENBANK	Sequence 5355	AF070548	GENBANK
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Sequence 5306	AF020038	GENBANK	Sequence 5358	AF070562	GENBANK
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Sequence 5307	AF020391 AF020768	GENBANK	Sequence 5360	AF070590 AF070652	GENBANK
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Sequence 5311	AF025998	GENBANK	Sequence 5363		GENBANK
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Sequence 5314	AF034209	GENBANK	Sequence 5366	AF078854	GENBANK
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Sequence 5319	AF038451	GENBANK	Sequence 5371	AF083190	GENBANK
Sequence 5320	AF038962	GENBANK	Sequence 5372	AF086168	GENBANK
Sequence 5321	AF039081	GENBANK	Sequence 5373	AF086245	GENBANK
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Sequence 5324	AF040958	GENBANK	Sequence 5376	AF089747	GENBANK
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Sequence 5330	AF047185	GENBANK	Sequence 5382	AF099149	GENBANK
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Sequence 5333	AF050641	GENBANK	Sequence 5385	AF102846	GENBANK
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Sequence 5336	AF052149	GENBANK	Sequence 5388	AF126181	GENBANK
Sequence 5337	AF052175	GENBANK	Sequence 5389	AF128528	GENBANK
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Sequence 5341	AF054838	GENBANK	Sequence 5393	AF147331	GENBANK
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Sequence 5345	AF060219	GENBANK	Sequence 5397	AF151800	GENBANK
Sequence 5346	AF061258	GENBANK	Sequence 5398	AF151840	GENBANK
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Sequence 5399	AF151846	GENBANK	•	D29640	GENBANK
Sequence 5400	AF151861	GENBANK	Sequence 5452 Sequence 5453	D29040 D29954	GENBANK
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Sequence 5402	AF151884	GENBANK	Sequence 5454		GENBANK
Sequence 5403	AF151893	GENBANK	Sequence 5455	D38037	
Sequence 5404	AF159056	GENBANK	Sequence 5456	D38549	GENBANK
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Sequence 5406	AF159295	GENBANK	Sequence 5458	D42044	GENBANK
Sequence 5407	AF168956	GENBANK	Sequence 5459	D42047	GENBANK
Sequence 5408	AF177291	GENBANK	Sequence 5460	D42138	GENBANK
Sequence 5409	AF177775	GENBANK	Sequence 5461	D43950	GENBANK
Sequence 5410	AF188298	GENBANK	Sequence 5462	D49372	GENBANK
Sequence 5411	AF188745	GENBANK	Sequence 5463	D49489	GENBANK
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Sequence 5413	AF190167	GENBANK	Sequence 5465	D50371	GENBANK
Sequence 5414	AJ010046	GENBANK	Sequence 5466	D50579	GENBANK
Sequence 5415	AJ010482	GENBANK	Sequence 5467	D50916	GENBANK
Sequence 5416	AJ012463	GENBANK	Sequence 5468	D50919	GENBANK
Sequence 5417	AJ012499	GENBANK	Sequence 5469	D55653	GENBANK
Sequence 5418	AJ130733	GENBANK	Sequence 5470	D63486	GENBANK
Sequence 5419	AJ223075	GENBANK	Sequence 5471	D64110	GENBANK
Sequence 5420	AJ223350	GENBANK	Sequence 5472	D80009	GENBANK
Sequence 5421	AJ223330 AJ224172	GENBANK	Sequence 5473	D82060	GENBANK
•	AJ249731	GENBANK	Sequence 5474	D83032	GENBANK
Sequence 5422	AL049705	GENBANK	Sequence 5475	D83735	GENBANK
Sequence 5423		GENBANK	Sequence 5476	D85181	GENBANK
Sequence 5424	AL049969		Sequence 5477	D86984	GENBANK
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Sequence 5426	AL050274	GENBANK	Sequence 5479	D87292	GENBANK
Sequence 5427	AL080089	GENBANK		D87453	GENBANK
Sequence 5428	AL080135	GENBANK	Sequence 5480	D87462	GENBANK
Sequence 5429	AL080209	GENBANK	Sequence 5481	D87666	GENBANK
Sequence 5430	AL096880	GENBANK	Sequence 5482		GENBANK
Sequence 5431	AL109672	GENBANK	Sequence 5483	D87684	
Sequence 5432	AL110183	GENBANK	Sequence 5484	D87735	GENBANK
Sequence 5433	AL117413	GENBANK	Sequence 5485	D87969	GENBANK
Sequence 5434	AL117550	GENBANK	Sequence 5486	D88674	GENBANK
Sequence 5435	AL117595	GENBANK	Sequence 5487	D90373	GENBANK
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Sequence 6385	AA747515	dbEST	Sequence 6437	AA923547	dbEST
Sequence 6386	AA748033	dbEST	Sequence 6438	AA926656	dbEST
Sequence 0300	~~i +0000	usaet i	2542555 5 700		

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Sequence 6439	AA928427	dbEST	Sequence 6491	AI085393	dbEST
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Sequence 6475 Sequence 6476	Al041847	dbEST	Sequence 6528	Al186545	dbEST
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Sequence 6544	Al217172	dbEST	Sequence 6596	Al349099	dbEST
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Sequence 6698	AI917110	dbEST	Sequence 6750	AW160696	dbEST
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Sequence 6751	AW161209	dbEST	Sequence 6803	N79712	dbEST
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Sequence 6761	D12135	dbEST	Sequence 6813	R24400	dbEST
Sequence 6762	D30921	dbEST	Sequence 6814	R25464	dbEST
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Sequence 6765	D82149	dbEST	Sequence 6817	R53758	dbEST
Sequence 6766	D82176	dbEST	Sequence 6818	R56382	dbEST
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Sequence 6769	F05508	dbEST	Sequence 6821	R70193	dbEST
Sequence 6770	H02725	dbEST	Sequence 6822	R70570	dbEST
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Sequence 6800	N62799	dbEST	Sequence 6852	Q97381	NUCPATENT
Sequence 6801	N63387	dbEST	Sequence 6853	T74203	NUCPATENT
Sequence 6802	N75468	dbEST	Sequence 6854	V07906	NUCPATENT
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Sequence 6857	V34226		Sequence 6908	AF052138	GENBANK
Sequence 6858	V34305	NUCPATENT NUCPATENT	Sequence 6909	AF054183	GENBANK
Sequence 6859	V35555	NUCPATENT	Sequence 6910	AF054987	GENBANK
Sequence 6860	V43604	NUCPATENT		AF055473	GENBANK
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Sequence 6864	V62427	NUCPATENT	Sequence 6914	AF068754	GENBANK
Sequence 6865	V69855	NUCPATENT	Sequence 6915		GENBANK
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Sequence 6867	V72125	NUCPATENT	Sequence 6917	AF070655	
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Sequence 6869	V84448	NUCPATENT	Sequence 6919	AF071748	GENBANK
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Sequence 7412	AI583257	dbEST	Sequence 7463	AL119319	dbEST
Sequence 7413	Al611738	dbEST	Sequence 7464	AW009700	dbEST
Sequence 7414	Al625204	dbEST	Sequence 7465	AW009886	dbEST

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Sequence 7485	AW183848	dbEST
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Sequence 7500	W73817	dbEST
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Sequence 1240

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Sequence 2802

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Sequence 2805

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Sequence 4712

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CATGATGGC

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Sequence 4716

Sequence 4717

Sequence 4718

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Sequence 4719

Sequence 4720

Sequence 4722

Sequence 4723

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Sequence 6889

Sequence 6890

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Sequence 6891

Sequence 6892

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Sequence 6893

Sequence 6894

Sequence 6895

Sequence 6896

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Sequence #	Acc No	Database hit	GI Number
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Sequence 7513	A26481	Genbank	1247452
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Sequence 7515	A29216	Genbank	1247528
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Sequence 7517	A36460	Genbank	2293778
Sequence 7518	A43521	Genbank	2298710
Sequence 7519	A58331	Genbank	3713993
Sequence 7520	A80239	Genbank	6092971
Sequence 7521	A81324	Genbank	6731650
Sequence 7522	A98295	Genbank	6781400
Sequence 7523	AB000360	Genbank	2547041
Sequence 7524	AB000468	Genbank	1843400
Sequence 7525	AB000516	Genbank	2723379
Sequence 7526	AB000516	Genbank	1813326
Sequence 7527	AB000624	Genbank	5748486
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Sequence 7530 Sequence 7531	AB001001	Genbank	2224528
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Sequence 7533	AB002299	Genbank	2224542
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Sequence 7536	AB002328	Genbank	2224600
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Sequence 7539 Sequence 7540	AB002340 AB002351	Genbank	2224646
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Sequence 7543	AB002365	Genbank	2224674
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Sequence 7547	AB002377	Genbank	6634024
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Sequence 7550	AB002387	Genbank	2224718
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Sequence 8305	AK001716	Genbank	7023150
Sequence 8306	AK001725	Genbank	7023165
Sequence 8307	AK001741	Genbank	7023192
Sequence 8308	AK001775	Genbank	7023255
	AK001770	Genbank	7023339
Sequence 8309		-	
Sequence 8310	AK001852	Genbank	7023376
Sequence 8311	AK001939	Genbank	7023513
Sequence 8312	AK001962	Genbank	7023551
Sequence 8313	AK002019	Genbank	7023647
Sequence 8314	AK002051	Genbank	7023698
Sequence 8315	AK002060	Genbank	7023713
	AK002000 AK002094	Genbank	7023767
Sequence 8316			
Sequence 8317	AK002136	Genbank	7023830
Sequence 8318	AK002165	Genbank	7023876
Sequence 8319	AK002212	Genbank	7023953
Sequence 8320	AL008639	Genbank	6599068
Sequence 8321	AL008721	Genbank	3171883
Sequence 8322	AL020995	Genbank	5881341
	AL021453	Genbank	3413288
Sequence 8323			2826890
Sequence 8324	AL021546	Genbank	
Sequence 8325	AL021707	Genbank	4582132
Sequence 8326	AL021807	Genbank	9367202
Sequence 8327	AL021917	Genbank	9367203
Sequence 8328	AL021978	Genbank	3135963
Sequence 8329	AL022157	Genbank	3319676
Sequence 8330	AL022313	Genbank	4200326
Sequence 8331	AL022318	Genbank	4826439
		Genbank	3355600
Sequence 8332	AL022339		
Sequence 8333	AL022394	Genbank	9581758
Sequence 8334	AL023803	Genbank	9408726
Sequence 8335	AL024498	Genbank	5924006
Sequence 8336	AL031387	Genbank	4493578
Sequence 8337	AL031427	Genbank	4835258
Sequence 8338	AL031680	Genbank	9368417
	AL031685	Genbank	9368423
Sequence 8339			
Sequence 8340	AL031733	Genbank	4826495
Sequence 8341	AL031778	Genbank	4153958
Sequence 8342	AL034343	Genbank	5931807
Sequence 8343	AL034379	Genbank	5918013
Sequence 8344	AL034380	Genbank	7981257
Sequence 8345	AL035420	Genbank	5918161
Sequence 8346	AL035551	Genbank	4826429
	AL035587	Genbank	6002306
Sequence 8347			
Sequence 8348	AL035659	Genbank	6562059
Sequence 8349	AL035683	Genbank	7288039
Sequence 8350	AL049219	Genbank	4499949

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Sequence 8351	AL049367	Genbank	4500158
Sequence 8352	AL049539	Genbank	6456855
Sequence 8353	AL049546	Genbank	4678775
Sequence 8354	AL049589	Genbank	5679567
Sequence 8355	AL049611	Genbank	5881343
Sequence 8356	AL049646	Genbank	9368462
•			
Sequence 8357	AL049712	Genbank	5629919
Sequence 8358	AL049713	Genbank	6006528
Sequence 8359	AL049742	Genbank	5419783
Sequence 8360	AL049795	Genbank	6010175
Sequence 8361	AL049801	Genbank	4902677
Sequence 8362	AL049821	Genbank	5791513
Sequence 8363	AL049943	Genbank	4884187
Sequence 8364	AL050011	Genbank	4884080
Sequence 8365	AL050028	Genbank	4884267
Sequence 8366	AL050107	Genbank	4884135
Sequence 8367	AL050118	Genbank	4884143
Sequence 8368	AL050135	Genbank	4884344
Sequence 8369	AL050187	Genbank	4884402
Sequence 8370	AL050208	Genbank	4884447
Sequence 8371	AL050208 AL050274		
•		Genbank	4886452
Sequence 8372	AL050332	Genbank	6010176
Sequence 8373	AL050367	Genbank	4914600
Sequence 8374	AL050396	Genbank	4914618
Sequence 8375	AL078581	Genbank	6967288
Sequence 8376	AL078594	Genbank	7161750
Sequence 8377	AL078611	Genbank	5019312
Sequence 8378	AL078621	Genbank	6013067
Sequence 8379	AL079298	Genbank	5102736
Sequence 8380	AL079352	Genbank	7799779
Sequence 8381	AL080080	Genbank	5262491
Sequence 8382	AL080155	Genbank	5262612
Sequence 8383	AL080157	Genbank	5262616
Sequence 8384	AL080172	Genbank	5262642
Sequence 8385	AL096738	Genbank	5419870
Sequence 8386	AL096740	Genbank	5419896
Sequence 8387	AL096752	Genbank	5419888
Sequence 8388	AL096791		
		Genbank	6066332
Sequence 8389	AL109788	Genbank	5725476
Sequence 8390	AL109794	Genbank	5725481
Sequence 8391	AL109925	Genbank	7105935
Sequence 8392	AL110244	Genbank	5817191
Sequence 8393	AL110268	Genbank	5817041
Sequence 8394	AL110273	Genbank	5817091
Sequence 8395	AL117398	Genbank	5911918
Sequence 8396	AL117409	Genbank	5912046
Sequence 8397	AL117429	Genbank	5911863
Sequence 8398	AL117482	Genbank	5911960
Sequence 8399	AL117500	Genbank	5912005
Sequence 8400	AL117572	Genbank	5912124
Sequence 8401	AL117573	Genbank	5912125
Sequence 8402	AL117579	Genbank	5912135
Sequence 8403	AL117594	Genbank	5912158
Sequence 8404	AL117607	Genbank	5912177
Sequence 8405	AL117694	Genbank	8248721
Sequence 8406	AL121575	Genbank	
Ocquence 0400	AL 12 13/3	Genbank	7264053

Sequence 8407	AL121790	Genbank	8919824
Sequence 8408	AL122020	Genbank	8176920
Sequence 8409	AL133036	Genbank	6453441
Sequence 8410	AL133064	Genbank	6453490
Sequence 8411	AL133078	Genbank	6453522
Sequence 8412	AL133215	Genbank	7228177
	AL133213 AL133246	Genbank	7594587
Sequence 8413			6706037
Sequence 8414	AL133353	Genbank	7019734
Sequence 8415	AL133455	Genbank	
Sequence 8416	AL133555	Genbank	6599122
Sequence 8417	AL133583	Genbank	6599167
Sequence 8418	AL133589	Genbank	6599176
Sequence 8419	AL133609	Genbank	6599219
Sequence 8420	AL135744	Genbank	6682291
Sequence 8421	AL135858	Genbank	7159622
Sequence 8422	AL136295	Genbank	6850939
Sequence 8423	AL136543	Genbank	6807646
Sequence 8424	AL137263	Genbank	6807692
Sequence 8425	AL137304	Genbank	6807768
Sequence 8426	AL137327	Genbank	6807818
Sequence 8427	AL137406	Genbank	6807955
Sequence 8428	AL137471	Genbank	6808079
Sequence 8429	AL137540	Genbank	6808218
	AL137548	Genbank	6808231
Sequence 8430	AL137546 AL137585	Genbank	6808304
Sequence 8431	AL137563 AL137641	Genbank	6808443
Sequence 8432		Genbank	6807931
Sequence 8433	AL137681	Genbank	6807935
Sequence 8434	AL137683	Genbank	6808256
Sequence 8435	AL137727	= :	6808323
Sequence 8436	AL137743	Genbank	6969146
Sequence 8437	AL139165	Genbank	4666256
Sequence 8438	AP000010	Genbank	3132356
Sequence 8439	AP000046	Genbank	
Sequence 8440	AP000066	Genbank	4579987
Sequence 8441	AP000493	Genbank	5926660
Sequence 8442	AP000501	Genbank	5926688
Sequence 8443	AP000529	Genbank	5931507
Sequence 8444	AP000694	Genbank	6693626
Sequence 8445	AP001115	Genbank	6899854
Sequence 8446	AP001137	Genbank	6970361
Sequence 8447	D00591	Genbank	220051
Sequence 8448	D01038	Genbank	220140
Sequence 8449	D10040	Genbank	219899
Sequence 8450	D10495	Genbank	520586
Sequence 8451	D12981	Genbank	219979
Sequence 8452	D13119	Genbank	285909
Sequence 8453	D13286	Genbank	496368
Sequence 8454	D13540	Genbank	220071
Sequence 8455	D13627	Genbank	286010
Sequence 8456	D13635	Genbank	285982
Sequence 8457	D13748	Genbank	219402
Sequence 8458	D14662	Genbank	285948
Sequence 8459	D14689	Genbank	285956
Sequence 8460	D14694	Genbank	603801
Sequence 8461	D14811	Genbank	285966
Sequence 8462	D16307	Genbank	303594
Sequence 0702	D .000.		

Table 3-3

Sequence 8463	D16562	Genbank	506336
Sequence 8464	D21063	Genbank	434752
Sequence 8465	D21163	Genbank	434758
-			
Sequence 8466	D21235	Genbank	498145
Sequence 8467	D21254	Genbank	575577
Sequence 8468	D21261	Genbank	434762
Sequence 8469	D21262	Genbank	434764
Sequence 8470	D21852	Genbank	434768
Sequence 8471	D25215	Genbank	517114
Sequence 8472	D26156	Genbank	505087
•			
Sequence 8473	D26181	Genbank	473712
Sequence 8474	D26351	Genbank	450470
Sequence 8475	D26485	Genbank	468934
Sequence 8476	D28473	Genbank	551621
Sequence 8477	D28476	Genbank	460710
Sequence 8478	D28589	Genbank	460714
Sequence 8479	D29954	Genbank	473940
		Genbank	473948
Sequence 8480	D29958		
Sequence 8481	D30756	Genbank	488500
Sequence 8482	D31762	Genbank	498149
Sequence 8483	D31764	Genbank	498153
Sequence 8484	D31767	Genbank	505091
Sequence 8485	D31885	Genbank	505097
Sequence 8486	D31888	Genbank	506340
Sequence 8487	D37766	Genbank	2429078
Sequence 8488	D38047	Genbank	1037163
Sequence 8489	D38048	Genbank	1531532
Sequence 8490	D38073	Genbank	862331
Sequence 8491	D38522	Genbank	559331
Sequence 8492	D38524	Genbank	633070
Sequence 8493	D38549	Genbank	559702
Sequence 8494	D38553	Genbank	559714
Sequence 8495	D38555	Genbank	559716
Sequence 8496		Genbank	1304117
•	D38616		
Sequence 8497	D42047	Genbank	577306
Sequence 8498	D42055	Genbank	577312
Sequence 8499	D42085	Genbank	577316
Sequence 8500	D43682	Genbank	1060913
Sequence 8501	D43949	Genbank	603952
Sequence 8502	D43950	Genbank	603954
Sequence 8503	D43951	Genbank	603956
Sequence 8504	D45021	Genbank	624872
Sequence 8505	D45131	Genbank	1304103
Sequence 8506	D49489	Genbank	1136742
Sequence 8507	D49490	Genbank	1072306
Sequence 8508	D50369	Genbank	2605589
Sequence 8509	D50372	Genbank	2605593
Sequence 8510	D50923	Genbank	1469188
Sequence 8511	D50929	Genbank	1469200
Sequence 8512	D55716	Genbank	1255616
Sequence 8513	D63475	Genbank	1665724
Sequence 8514	D63478	Genbank	1469869
Sequence 8515	D63479	Genbank	6633998
Sequence 8516	D63486	Genbank	1469885
Sequence 8517	D63780	Genbank	2196444
Sequence 8518	D63861	Genbank	1769811

<u>Table 3-3</u>

			004444
Sequence 8519	D63875	Genbank	961441
Sequence 8520	D78014	Genbank	1330241
Sequence 8521	D79205	Genbank	1754620
Sequence 8522	D79984	Genbank	1136385
		Genbank	1136397
Sequence 8523	D79991		
Sequence 8524	D80000	Genbank	1136415
Sequence 8525	D80009	Genbank	1136433
Sequence 8526	D80010	Genbank	1136435
Sequence 8527	D82348	Genbank	1311461
Sequence 8528	D83077	Genbank	1304131
	D83778	Genbank	1228038
Sequence 8529		Genbank	1228044
Sequence 8530	D83781		
Sequence 8531	D83782	Genbank	1228046
Sequence 8532	D84064	Genbank	2618587
Sequence 8533	D86198	Genbank	3062805
Sequence 8534	D86962	Genbank	1503997
Sequence 8535	D86965	Genbank	1504003
Sequence 8536	D86977	Genbank	1504027
	D86978	Genbank	1504029
Sequence 8537			6634002
Sequence 8538	D86985	Genbank	
Sequence 8539	D86987	Genbank	3869186
Sequence 8540	D87127	Genbank	1817551
Sequence 8541	D87436	Genbank	1665766
Seguence 8542	D87447	Genbank	1665782
Sequence 8543	D87452	Genbank	1665792
Sequence 8544	D87464	Genbank	1665812
Sequence 8545	D87470	Genbank	1665822
	D87717	Genbank	1663709
Sequence 8546		Genbank	1620021
Sequence 8547	D87735		
Sequence 8548	D88674	Genbank	2641951
Sequence 8549	D88687	Genbank	1843433
Sequence 8550	D89980	Genbank	2804272
Sequence 8551	D90391	Genbank	219491
Sequence 8552	E01094	Genbank	2169353
Sequence 8553	E01787	Genbank	2170040
Sequence 8554	E01816	Genbank	2170068
Sequence 8555	E01956	Genbank	2170204
Sequence 8556	E02164	Genbank	2170402
	E02628	Genbank	2170856
Sequence 8557			2174903
Sequence 8558	E06721	Genbank	
Sequence 8559	E12458	Genbank	3251291
Sequence 8560	E13330	Genbank	3252135
Sequence 8561	E14580	Genbank	5709263
Sequence 8562	E14872	Genbank	5709555
Sequence 8563	E15239	Genbank	5709922
Sequence 8564	E15695	Genbank	5710378
Sequence 8565	E17361	Genbank	5712044
	J02642	Genbank	182862
Sequence 8566		Genbank	
Sequence 8567	J02923		189501
Sequence 8568	J03007	Genbank	179211
Sequence 8569	J03015	Genbank	337755
Sequence 8570	J03202	Genbank	186916
Sequence 8571	J03473	Genbank	337423
Sequence 8572	J03518	Genbank	340389
Sequence 8573	J03528	Genbank	188671
Sequence 8574	J03575	Genbank	189737
30400100 0014	555575	COIDGIII	

Sequence 8575	J03576	Genbank	189753
Sequence 8576	J03592	Genbank	339722
Sequence 8577	J03799	Genbank	186840
Sequence 8578	J03827	Genbank	340418
Sequence 8579	J04031	Genbank	187464
Sequence 8580	J04088	Genbank	292829
	J04060 J04164	Genbank	177801
Sequence 8581		Genbank	551173
Sequence 8582	J04173		
Sequence 8583	J04444	Genbank	181239
Sequence 8584	J04543	Genbank	338243
Sequence 8585	J04611	Genbank	178649
Sequence 8586	J04615	Genbank	338246
Sequence 8587	J04988	Genbank	184422
Sequence 8588	J05594	Genbank	1203981
Sequence 8589	J05633	Genbank	186504
Sequence 8590	K00558	Genbank	340020
Sequence 8591	L00160	Genbank	189904
Sequence 8592	L00635	Genbank	292032
Sequence 8593	L01457	Genbank	179282
Sequence 8594	L04733	Genbank	307084
Sequence 8595	L05093	Genbank	401844
Sequence 8596	L05425	Genbank	179284
Sequence 8597	L06237	Genbank	437000
Sequence 8598	L06328	Genbank	340200
Sequence 8599	L06419	Genbank	190073
Sequence 8600	L07044	Genbank	291894
Sequence 8601	L08238	Genbank	307198
Sequence 8602	L00250	Genbank	307374
Sequence 8603	L09561	Genbank	3192937
Sequence 8604	L11931	Genbank	307421
	L11931	Genbank	307423
Sequence 8605	L11932 L13850	Genbank	292165
Sequence 8606	L13030	Genbank	307437
Sequence 8607	L14070 L14283	Genbank	307355
Sequence 8608		Genbank	348238
Sequence 8609	L14599		
Sequence 8610	L14837	Genbank	292937
Sequence 8611	L16783	Genbank	452438
Sequence 8612	L19597	Genbank	306467
Sequence 8613	L19956	Genbank	306456
Sequence 8614	L20941	Genbank	507251
Sequence 8615	L22005	Genbank	388308
Sequence 8616	L22009	Genbank	347313
Sequence 8617	L22740	Genbank	409222
Sequence 8618	L23959	Genbank	414316
Sequence 8619	L25270	Genbank	457136
Sequence 8620	L25876	Genbank	414584
Sequence 8621	L25880	Genbank	450272
Sequence 8622	L26524	Genbank	457232
Sequence 8623	L27211	Genbank	558656
Sequence 8624	L27711	Genbank	808006
Sequence 8625	L29216	Genbank	632965
Sequence 8626	L29296	Genbank	559042
Sequence 8627	L29555	Genbank	522196
Sequence 8628	L34075	Genbank	508481
Sequence 8629	L36056	Genbank	561631
Sequence 8630	L36818	Genbank	556191

Sequence 8631	L38961	Genbank	624703
Sequence 8632	L39068	Genbank	994714
Sequence 8633	L39210	Genbank	1702963
Sequence 8634	L39211	Genbank	755645
Sequence 8635	L40326	Genbank	695361
Sequence 8636	L40586	Genbank	1222629
Sequence 8637	L40933	Genbank	1160964
Sequence 8638	L41607	Genbank	886293
Sequence 8639	L41816	Genbank	790789
Sequence 8640	L42379	Genbank	1203964
Sequence 8641	L42542	Genbank	974142
	L42809	Genbank	1160966
Sequence 8642	L42609 L43509	Genbank	896282
Sequence 8643	L43309 L47162	Genbank	1082035
Sequence 8644	L47162 L47647	Genbank	1002033
Sequence 8645			1405422
Sequence 8646	L49345	Genbank Genbank	1196416
Sequence 8647	L54057		1280205
Sequence 8648	L77701	Genbank	1479980
Sequence 8649	L78464	Genbank	
Sequence 8650	L78669	Genbank	1374862
Sequence 8651	M11147	Genbank	182513
Sequence 8652	M11560	Genbank	178350
Sequence 8653	M12938	Genbank	182515
Sequence 8654	M13520	Genbank	179459
Sequence 8655	M13755	Genbank	184570
Sequence 8656	M14326	Genbank	340005
Sequence 8657	M14631	Genbank	183416
Sequence 8658	M15182	Genbank	183232
Sequence 8659	M16038	Genbank	187268
Sequence 8660	M16247	Genbank	178044
Sequence 8661	M16279	Genbank	188542
Sequence 8662	M17081	Genbank	178151
Sequence 8663	M17851	Genbank	182860
Sequence 8664	M17885	Genbank	190231
Sequence 8665	M17886	Genbank	190233
Sequence 8666	M18112	Genbank	190166
Sequence 8667	M18157	Genbank	186640
Sequence 8668	M18366	Genbank	179131
Sequence 8669	M19961	Genbank	180940
Sequence 8670	M20372	Genbank	189372
Sequence 8671	M20776	Genbank	180909
Sequence 8672	M21154	Genbank	178517
Sequence 8673	M21575	Genbank	1311702
Sequence 8674	M21895	Genbank	189523
Sequence 8675	M22382	Genbank	190126
Sequence 8676	M22538	Genbank	986883
Sequence 8677	M22803	Genbank	190379
Sequence 8678	M22919	Genbank	189016
Sequence 8679	M22920	Genbank	189021
Sequence 8680	M22960	Genbank	190282
Sequence 8681	M23254	Genbank	511636
Sequence 8682	M23419	Genbank	181996
Sequence 8683	M23725	Genbank	189997
Sequence 8684	M24194	Genbank	187701
Sequence 8685	M24355	Genbank	182604
Sequence 8686	M24486	Genbank	190785
Dequence 0000	14127700	Combanit	,

Sequence 8687	M24543	Genbank	341200
Sequence 8688	M26066	Genbank	189854
Sequence 8689	M26325	Genbank	186688
Sequence 8690	M26663	Genbank	618463
Sequence 8691	M29366	Genbank	181979
•	M29551	Genbank	180708
Sequence 8692			
Sequence 8693	M31159	Genbank	183115
Sequence 8694	M31511	Genbank	179907
Sequence 8695	M31606	Genbank	189940
Sequence 8696	M33308	Genbank	340236
Sequence 8697	M34055	Genbank	190791
Sequence 8698	M34064	Genbank	416292
Sequence 8699	M34081	Genbank	337458
Sequence 8700	M34088	Genbank	182122
Sequence 8701	M34668	Genbank	190738
Sequence 8702	M37499	Genbank	188886
Sequence 8703	M38188	Genbank	189378
Sequence 8704	M55153	Genbank	339520
•		Genbank	184483
Sequence 8705	M55618	•••••	
Sequence 8706	M57567	Genbank	178986
Sequence 8707	M58028	Genbank	340071
Sequence 8708	M59371	Genbank	181943
Sequence 8709	M59849	Genbank	182591
Sequence 8710	M61715	Genbank	340367
Sequence 8711	M62810	Genbank	188563
Sequence 8712	M64098	Genbank	183891
Sequence 8713	M64571	Genbank	187382
Sequence 8714	M64788	Genbank	190855
Sequence 8715	M65217	Genbank	184404
Sequence 8716	M67480	Genbank	190364
Sequence 8717	M69066	Genbank	188625
Sequence 8718	M69181	Genbank	641957
Sequence 8719	M73816	Genbank	183609
	M74491	Genbank	178161
Sequence 8720		Genbank	184020
Sequence 8721	M75126		
Sequence 8722	M76729	Genbank	189519
Sequence 8723	M80254	Genbank	181273
Sequence 8724	M80335	Genbank	189966
Sequence 8725	M80646	Genbank	338701
Sequence 8726	M80776	Genbank	179334
Sequence 8727	M81934	Genbank	180172
Sequence 8728	M83651	Genbank	431032
Sequence 8729	M84326	Genbank	178163
Sequence 8730	M84739	Genbank	179881
Sequence 8731	M85038	Genbank	190189
Sequence 8732	M86492	Genbank	183369
Sequence 8733	M91670	Genbank	181915
Sequence 8734	M95627	Genbank	870802
Sequence 8735	M96684	Genbank	190749
	M96803	Genbank	338442
Sequence 8736			
Sequence 8737	M96995	Genbank	181975
Sequence 8738	M97501	Genbank	180621
Sequence 8739	M97934	Genbank	186566
Sequence 8740	M97935	Genbank	2281070
Sequence 8741	M98478	Genbank	339577
Sequence 8742	M99422	Genbank	183662

Sequence 8743	S60710	Genbank	300301
Sequence 8744	S62138	Genbank	386158
Sequence 8745	S67779	Genbank	459372
Sequence 8746	S69272	Genbank	546087
Sequence 8747	S70114	Genbank	546696
Sequence 8748	S73498	Genbank	688010
Sequence 8749	S74678	Genbank	241477
Sequence 8750	S74799	Genbank	797365
Sequence 8751	S75174	Genbank	802120
Sequence 8752	S82081	Genbank	1488412
Sequence 8753	U00930	Genbank	405043
Sequence 8754	U00947	Genbank	405049
Sequence 8755	U02390	Genbank	409928
Sequence 8756	U02493	Genbank	407307
Sequence 8757	U03886	Genbank	458225
	U05572	Genbank	1419373
Sequence 8758		Genbank	514264
Sequence 8759	U07563		706838
Sequence 8760	U07681	Genbank	469038
Sequence 8761	U07810	Genbank	
Sequence 8762	U07919	Genbank	995897
Sequence 8763	U08377	Genbank	508230
Sequence 8764	U09220	Genbank	799338
Sequence 8765	U09284	Genbank	516011
Sequence 8766	U09716	Genbank	606827
Sequence 8767	U09813	Genbank	1008454
Sequence 8768	U10248	Genbank	984280
Sequence 8769	U10323	Genbank	532312
Sequence 8770	U10439	Genbank	577169
Sequence 8771	U10691	Genbank	533522
Sequence 8772	U10886	Genbank	1685074
Sequence 8773	U12424	Genbank	525319
Sequence 8774	U12535	Genbank	530822
Sequence 8775	U13022	Genbank	537293
Sequence 8776	U13991	Genbank	562076
Sequence 8777	U14193	Genbank	555899
Sequence 8778	U14966	Genbank	550012
Sequence 8779	U14969	Genbank	550018
Sequence 8780	U14970	Genbank	550020
Sequence 8781	U14971	Genbank	550022
Sequence 8782	U15158	Genbank	576622
Sequence 8783	U16660	Genbank	564064
Sequence 8784	U18121	Genbank	915283
Sequence 8785	U18197	Genbank	603073
Sequence 8786	U18937	Genbank	899108
Sequence 8787	U19145	Genbank	914904
Sequence 8788	U19252	Genbank	808035
Sequence 8789	U20476	Genbank	3077819
Sequence 8790	U21910	Genbank	736402
Sequence 8791	U22055	Genbank	799176
Sequence 8792	U22815	Genbank	930340
Sequence 8793	U23028	Genbank	806853
Sequence 8794	U24169	Genbank	1215668
Sequence 8795	U24223	Genbank	1215670
	U25816	Genbank	837262
Sequence 8796	U26425	Genbank	836664
Sequence 8797	U27515	Genbank	4096274
Sequence 8798	027010	Genbank	7030214

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Sequence 9328	AA402063	EST	2056854
Sequence 9329	AA403047	EST	2055609
Sequence 9330	AA406425	EST	2064410
Sequence 9331	AA412426	EST	2071129
Sequence 9332	AA417300	EST	2077408
	AA417999	EST	2079818
Sequence 9333			
Sequence 9334	AA418045	EST	2079864
Sequence 9335	AA418096	EST	2079897
Sequence 9336	AA418249	EST	2080078
Sequence 9337	AA421643	EST	2100459
Sequence 9338	AA424317	EST	2103287
Sequence 9339	AA425387	EST	2106170
Sequence 9340	AA425619	EST	2106375
Sequence 9341	AA425658	EST	2106433
	AA425751	EST	2106453
Sequence 9342			
Sequence 9343	AA425761	EST	2106481
Sequence 9344	AA427920	EST	2112054
Sequence 9345	AA428948	EST	2110491
Sequence 9346	AA429871	EST	2113108
Sequence 9347	AA431577	EST	2115285
Sequence 9348	AA433865	EST	2138779
Sequence 9349	AA436966	EST	2141880
Sequence 9350	AA442297	EST	2154175
Sequence 9351	AA442687	EST	2154565
Sequence 9352	AA446391	EST	2159056
Sequence 9353	AA448292	EST	2161962
Sequence 9354	AA449483	EST	2163233
Sequence 9355	AA449501	EST	2163251
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Sequence 9358	AA450166	EST	2163916
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Sequence 9359	AA450276	EST	2164026
Sequence 9360	AA452716	EST	2166385
Sequence 9361	AA453565	EST	2167234
Sequence 9362	AA453899	EST	2167568
Sequence 9363	AA456454	EST	2179030
Sequence 9364	AA457365	EST	2180085
Sequence 9365	AA458613	EST	
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Sequence 9366	AA459167	EST	2184074
Sequence 9367	AA461383	EST	2186503
Sequence 9368	AA463367	EST	2188251
Sequence 9369	AA465402	EST	2191569
Sequence 9370	AA469033	EST	2195567
Sequence 9371	AA470067	EST	2197376
Sequence 9372	AA476876	EST	2205087
Sequence 9373	AA478162	EST	2206796
Sequence 9374	AA479499	EST	2208055
Sequence 9375	AA480816	EST	2210368
Sequence 9376	AA481233	EST	2210785
Sequence 9377	AA481838	EST	2209516
Sequence 9378	AA485255	EST	2214474
Sequence 9379	AA485520	EST	2214739
Sequence 9380	AA488447	EST	2215878
Sequence 9381	AA488488	EST	2215919
Sequence 9382	AA488513	EST	2215944
Sequence 9383	AA489117	EST	2218719
Sequence 9384	AA489288	EST	2218890
Sequence 9385	AA490389	EST	2219562
Sequence 9386	AA491213	EST	2220386
Sequence 9387	AA491638	EST	2221200
Sequence 9388	AA494334	EST	2224121
Sequence 9389	AA494355	EST	2224121
Sequence 9390	AA494387	EST	2224174
Sequence 9391	AA495912	EST	2229233
Sequence 9392	AA496486	EST	2229233
	AA501442		
Sequence 9393		EST	2236409
Sequence 9394	AA504269	EST	2240429
Sequence 9395	AA505816	EST	2241953
Sequence 9396	AA507818	EST	2244257
Sequence 9397	AA508038	EST	2244477
Sequence 9398	AA514302	EST	2253810
Sequence 9399	AA516531	EST	2256055
Sequence 9400	AA522888	EST	2263600
Sequence 9401	AA527374	EST	2269443
Sequence 9402	AA532530	EST	2276784
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Sequence 9406	AA535663	EST	2279916
Sequence 9407	AA535845	EST	2280098
Sequence 9408	AA552507	EST	2322761
Sequence 9409	AA558357	EST	2328834
Sequence 9410	AA565548	EST	2337187
Sequence 9411	AA565867	EST	2337506
Sequence 9412	AA568785	EST	2341839
Sequence 9413	AA573427	EST	2347955
Sequence 9414	AA574223	EST	2348738

Sequence 9415	AA576016	EST	2350531
Sequence 9416	AA576889	EST	2354363
Sequence 9417	AA579611	EST	2357795
		EST	2358957
Sequence 9418	AA581185		
Sequence 9419	AA587264	EST	2398078
Sequence 9420	AA587570	EST	2397245
Sequence 9421	AA594562	EST	2409912
Sequence 9422	AA598697	EST	2432369
Sequence 9423	AA600084	EST	2433709
Sequence 9424	AA600355	EST	2433980
Sequence 9425	AA601289	EST	2434914
Sequence 9426	AA603876	EST	2437737
Sequence 9427	AA604860	EST	2445724
Sequence 9428	AA608613	EST	2457041
	AA609417	EST	2457845
Sequence 9429			2463706
Sequence 9430	AA612668	EST	
Sequence 9431	AA625213	EST	2537598
Sequence 9432	AA626629	EST	2539016
Sequence 9433	AA627320	EST	2540337
Sequence 9434	AA628491	EST	2540878
Sequence 9435	AA628520	EST	2540907
Sequence 9436	AA631993	EST	2555407
Sequence 9437	AA634958	EST	2558172
Sequence 9438	AA640716	EST	2565966
Sequence 9439	AA641939	EST	2567157
Sequence 9440	AA652388	EST	2584040
		EST	2595847
Sequence 9441	AA659693		
Sequence 9442	AA662024	EST	2616115
Sequence 9443	AA663649	EST	2617640
Sequence 9444	AA668886	EST	2630385
Sequence 9445	AA679400	EST	2659922
Sequence 9446	AA679621	EST	2660143
Sequence 9447	AA687221	EST	2675412
Sequence 9448	AA687495	EST	2675686
Sequence 9449	AA693932	EST	2694870
Sequence 9450	AA694188	EST	2695126
Sequence 9451	AA700208	EST	2703171
Sequence 9452	AA701138	EST	2704303
Sequence 9453	AA704533	EST	2714451
	AA705937	EST	2715855
Sequence 9454			2718920
Sequence 9455	AA709002	EST	
Sequence 9456	AA716423	EST	2728697
Sequence 9457	AA722455	EST	2740162
Sequence 9458	AA722786	EST	2740493
Sequence 9459	AA730400	EST	2751604
Sequence 9460	AA731394	EST	2753550
Sequence 9461	AA742260	EST	2784260
Sequence 9462	AA746310	EST	2786296
Sequence 9463	AA746748	EST	2786734
Sequence 9464	AA747197	EST	2787155
Sequence 9465	AA749387	EST	2789345
Sequence 9466	AA757433	EST	2805296
	AA761101	EST	2810031
Sequence 9467			2817145
Sequence 9468	AA765907	EST	
Sequence 9469	AA769588	EST	2820826
Sequence 9470	AA772570	EST	2824353

<u>Table 3-3</u>

Sequence 9471	AA774779	EST	2834113
Sequence 9472	AA775625	EST	2834959
	AA805474	EST	2874224
Sequence 9473			
Sequence 9474	AA808205	EST	2877611
Sequence 9475	AA808239	EST	2877645
Sequence 9476	AA813991	EST	2883587
Sequence 9477	AA814182	EST	2883778
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Sequence 9478	AA814817	EST	2884413
Sequence 9479	AA826906	EST	2900903
Sequence 9480	AA830072	EST	2903171
Sequence 9481	AA831991	EST	2905090
Sequence 9482	AA833749	EST	2908517
Sequence 9483	AA837407	EST	2912606
Sequence 9484	AA838157	EST	2913814
Sequence 9485	AA843207	EST	2929725
Sequence 9486	AA843238	EST	2929756
Sequence 9487	AA843878	EST	2930329
Sequence 9488	AA843908	EST	2930359
Sequence 9489	AA857964	EST	2946266
Sequence 9490	AA864964	EST	2959277
Sequence 9491	AA868452	EST	2963897
Sequence 9492	AA868722	EST	2964167
Sequence 9493	AA872507	EST	2968685
Sequence 9494	AA872913	EST	2969035
Sequence 9495	AA872991	EST	2969113
	AA877350		
Sequence 9496		EST	2986427
Sequence 9497	AA877789	EST	2986754
Sequence 9498	AA902103	EST	3037293
Sequence 9499	AA902974	EST	3038097
Sequence 9500	AA903253	EST	3038376
Sequence 9501	AA905330	EST	3040453
Sequence 9502	AA906868	EST	3042112
•			3042646
Sequence 9503	AA907186	EST	
Sequence 9504	AA908460	EST	3047865
Sequence 9505	AA913203	EST	3052595
Sequence 9506	AA916003	EST	3055395
Sequence 9507	AA927612	EST	3077033
Sequence 9508	AA931566	EST	3085952
Sequence 9509	AA932278	EST	3086516
Sequence 9510	AA968524	EST	3143704
Sequence 9511	AA973790	EST	3148970
Sequence 9512	AA976114	EST	3151906
Sequence 9513	AA976653	EST	3154099
Sequence 9514	AA984122	EST	3162647
Sequence 9515	AA988106	EST	3173470
	AA988525		3174217
Sequence 9516		EST	
Sequence 9517	AA988576	EST	3174268
Sequence 9518	AA991580	EST	3178069
Sequence 9519	AF001542	EST	2529714
Sequence 9520	AI015791	EST	3230127
Sequence 9521	AI017095	EST	3231431
Sequence 9522	AI017902	EST	3232238
Sequence 9523	AI017902 AI018625		
		EST	3233144
Sequence 9524	Al022209	EST	3239562
Sequence 9525	AI027035	EST	3244551
Sequence 9526	AI033991	EST	3254944

<u>Table 3-3</u>

Sequence 9527	AI034371	EST	3255324
Sequence 9528		EST	3279796
Sequence 9529		EST	3307000
		EST	3307991
Sequence 9530			
Sequence 9531	AI056337	EST	3330203
Sequence 9532		EST	3330268
Sequence 9533	AI061175	EST	3336543
Sequence 9534	AI064910	EST	6359182
Sequence 9535		EST	3367285
Sequence 9536		EST	3405808
Sequence 9537		EST	3404782
Sequence 9538		EST	3412625
		EST	3415761
Sequence 9539		EST	3416748
Sequence 9540			3418572
Sequence 9541		EST	
Sequence 9542		EST	3423217
Sequence 9543		EST	3423339
Sequence 9544	Al085616	EST	3424039
Sequence 9545		EST	3427956
Sequence 9546		EST	3429256
Sequence 9547		EST	3430232
Sequence 9548		EST	3433490
Sequence 9549		EST	3434426
		EST	6360037
Sequence 9550		EST	3539184
Sequence 9551			
Sequence 9552		EST	3593306
Sequence 9553		EST	3594519
Sequence 9554		EST	3597764
Sequence 9555	5 Al133074	EST	6360390
Sequence 9556		EST	6360404
Sequence 9557		EST	6360590
Sequence 9558		EST	6360742
Sequence 9559		EST	3645736
Sequence 9560		EST	3658773
Sequence 9561		EST	3674515
		EST	3675788
Sequence 9562		EST	3678899
Sequence 9563			
Sequence 9564		EST	3679615
Sequence 9565		EST	3693133
Sequence 9566		EST	3694377
Sequence 9567	7 Al168316	EST	3701486
Sequence 9568	3 Al168348	EST	3701518
Sequence 9569		EST	3737091
Sequence 9570		EST	3737795
Sequence 957		EST	3740684
Sequence 9572		EST	3744581
Sequence 9573		EST	3750824
		EST	3751797
Sequence 9574			3751800
Sequence 957		EST	
Sequence 9576		EST	3754743
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Sequence 9578		EST	6361479
Sequence 9579	9 Al207868	EST	3769810
Sequence 9580		EST	3777649
Sequence 958		EST	3778054
Sequence 9582		EST	3785139
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Sequence 9583	Al217246	EST	3797061
Sequence 9584	AI244268	EST	3839665
Sequence 9585	Al248386	EST	3843783
Sequence 9586	Al249242	EST	3845771
	Al251739	EST	3848268
Sequence 9587		EST	3870183
Sequence 9588	Al261980		
Sequence 9589	Al262251	EST	3870454
Sequence 9590	Al267162	EST	3886329
Sequence 9591	Al267848	EST	3887015
Sequence 9592	AI268335	EST	3887502
Sequence 9593	AI268362	EST	3887529
Sequence 9594	AI270685	EST	3889852
Sequence 9595	AI275466	EST	3897740
Sequence 9596	Al276933	EST	3899201
Sequence 9597	Al277443	EST	3899711
Sequence 9598	Al278328	EST	3900596
Sequence 9599	Al279625	EST	3917859
Sequence 9600	Al280561	EST	3918794
Sequence 9601	Al281394	EST	3919627
Sequence 9602	Al283373	EST	3921606
Sequence 9603	Al283578	EST	3921811
	Al284482	EST	3922715
Sequence 9604	Al290052	EST	3931718
Sequence 9605		EST	3931777
Sequence 9606	Al290111	EST	3931820
Sequence 9607	Al201087	EST	3933861
Sequence 9608	Al291087	EST	3934868
Sequence 9609	Al292094		3958628
Sequence 9610	Al298974	EST	4069675
Sequence 9611	Al333116	EST	
Sequence 9612	Al334177	EST	4070736
Sequence 9613	Al334889	EST	4071816
Sequence 9614	Al337358	EST	4074285
Sequence 9615	Al339284	EST	4076211
Sequence 9616	Al341138	EST	4078065
Sequence 9617	Al341291	EST	4078218
Sequence 9618	Al342090	EST	4079017
Sequence 9619	Al343404	EST	4080610
Sequence 9620	Al344189	EST	4081395
Sequence 9621	Al348588	EST	4085794
Sequence 9622	Al349337	EST	4086543
Sequence 9623	Al349957	EST	4087163
Sequence 9624	Al350524	EST	4087730
Sequence 9625	Al352184	EST	4089390
Sequence 9626	Al355277	EST	4095430
Sequence 9627		EST	4108082
Sequence 9628		EST	4109187
Sequence 9629		EST	4109203
Sequence 9630		EST	4112302
Sequence 9631	Al361796	EST	4113417
Sequence 9632		EST	4153694
Sequence 9633		EST	4188753
Sequence 9634		EST	4189502
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Sequence 9637		EST	4222816
Sequence 9638		EST	4266517
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Sequence 9639	AI435112	EST	4300898
		EST	4304204
Sequence 9640			
Sequence 964	1 Al439588	EST	4305374
Sequence 9642	2 Al445716	EST	4290157
Sequence 9643		EST	4307708
		EST	4310301
Sequence 9644			
Sequence 964		EST	4310833
Sequence 9646	3 Al458599	EST	4311178
Sequence 9647		EST	4311337
Sequence 9648		EST	4313101
		EST	4330509
Sequence 9649			
Sequence 9650		EST	4334174
Sequence 965	1 Al472566	EST	4325611
Sequence 9652		EST	4372243
Sequence 9653		EST	4373422
		EST	4393035
Sequence 9654			
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Sequence 9656	3 Al525541	EST	4439676
Sequence 965	7 Al525654	EST	4439789
Sequence 9658		EST	4452977
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Sequence 9659			4511314
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Sequence 966		EST	4522378
Sequence 966	2 Al564719	EST	4523176
Sequence 966		EST	4533989
Sequence 966		EST	4564628
Sequence 966		EST	4564712
		EST	4565474
Sequence 966			
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Sequence 966	8 Al608796	EST	4617963
Sequence 966	9 Al609087	EST	4618254
Sequence 967	0 Al610880	EST	4620047
Sequence 967		EST	4648915
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		EST	4665154
Sequence 967			
Sequence 967		EST	4665217
Sequence 967	5 Al632470	EST	4683800
Sequence 967	6 Al633000	EST	4684330
Sequence 967		EST	4690060
Sequence 967		EST	4690795
		EST	4703716
Sequence 967			
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Sequence 968		EST	4735856
Sequence 968	2 Al672492	EST	4852223
Sequence 968	3 Al684595	EST	4895889
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Sequence 968		EST	4901383
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Sequence 969	4 AI750105	EST	5128369

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Sequence 9696	Al751459	EST	5129723
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Sequence 9699	AI752414	EST	5130678
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Sequence 9702	AI761315	EST	5176982
Sequence 9703	AI765501	EST	5232010
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Sequence 9710	AI800215	EST	5365687
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Sequence 9712	AI807798	EST	5394364
Sequence 9713	Al808865	EST	5395431
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	Al814463	EST	5425678
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Sequence 9716	AI814513	EST	5431038
Sequence 9717	AI815492		5431984
Sequence 9718	AI816438	EST	
Sequence 9719	AI818683	EST	5437762
Sequence 9720	AI819405	EST	5438484
Sequence 9721	AI828075	EST	5448746 5537480
Sequence 9722	AI863382	EST	5527489 5547050
Sequence 9723	AI873001	EST	5547050 5547550
Sequence 9724	AI873501	EST	5547550 5548375
Sequence 9725	Al874326	EST	5548375
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Sequence 9727	Al884783	EST	5589947
Sequence 9728	AI885021	EST	5590185
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Sequence 9730	Al888327	EST	5593577
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Sequence 9733	Al904447	EST	6494834
Sequence 9734	Al906328	EST	6496715
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Sequence 9737	Al918605	EST	5638460
Sequence 9738	Al924534	EST	5660498
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Sequence 9742	Al950991	EST	5743301
Sequence 9743	Al970095	EST	5766921
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Sequence 9745	Al982834	EST	5810053
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Sequence 9747	AL035972	EST	5405602
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Sequence 9749	AL036415	EST	5405971
Sequence 9750	AL037386	EST	5406786
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Sequence 9752	AL039569	EST	5408608
Sequence 9753	AL039961	EST	5408942
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Sequence 9756	AL041673	EST	5421022
Sequence 9757	AL042613	EST	5935505
Sequence 9758	AL042680	EST	5422129
Sequence 9759	AL042725	EST	5422174
	AL042986	EST	5935655
Sequence 9760			
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Sequence 9762	AL043542	EST	5422929
Sequence 9763	AL043672	EST	5423059
Sequence 9764	AL045757	EST	5936079
Sequence 9765	AL045793	EST	5433905
Sequence 9766	AL045936	EST	5434039
Sequence 9767	AL045937	EST	5936123
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Sequence 9768	AL045976	EST	5434076
Sequence 9769	AL046015	EST	5434109
Sequence 9770	AL047001	EST	5435057
Sequence 9771	AL047332	EST	4727279
Sequence 9772	AL047890	EST	4728078
Sequence 9773	AL048554	EST	5936565
Sequence 9774	AL110421	EST	5866029
Sequence 9775	AL118756	EST	5924655
Sequence 9776	AL118845	EST	5924744
Sequence 9777	AL119809	EST	5925708
Sequence 9778	AL120711	EST	5926610
Sequence 9779	AL120865	EST	5926866
Sequence 9780	AL120911	EST	5926912
Sequence 9781	AL121473	EST	5927474
Sequence 9782	AL121558	EST	5927559
Sequence 9783	AL133779	EST	6601967
	AL133900		
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Sequence 9786	AL134795	EST	6602982
Sequence 9787	AL134938	EST	6603125
Sequence 9788	AL135007	EST	6603194
Sequence 9789	AL135023	EST	6603210
Sequence 9790	AL135290	EST	6603477
Sequence 9791	AL135343	EST	6603530
Sequence 9792	AL135362	EST	6603549
	AL135502 AL135502	EST	6603689
Sequence 9793			
Sequence 9794	AL135600	EST	6603787
Sequence 9795	AL138042	EST	6854722
Sequence 9796	AW002442	EST	5849358
Sequence 9797	AW014499	EST	5863256
Sequence 9798	AW014727	EST	5863484
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Sequence 9802	AW062837	EST	6014222
Sequence 9803	AW068104	EST	6023102
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Sequence 9805	AW070690	EST	6025688
Sequence 9806	AW080672	EST	6035905
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Sequence 9808	AW118447	EST	6087031
Sequence 9809	AW148635	EST	6196531
Sequence 9810	AW149925	EST	6197821
		EST	6228632
Sequence 9811	AW157231		
Sequence 9812	AW160468	EST	6299501
Sequence 9813	AW160833	EST	6299866
Sequence 9814	AW161807	EST	6300840
Sequence 9815	AW161978	EST	6301011
Sequence 9816	AW162587	EST	6301620
Sequence 9817	AW162761	EST	6301794
Sequence 9818	AW162774	EST	6301807
Sequence 9819	AW163485	EST	6302518
Sequence 9820	AW163783	EST	6302816
•	AW175627	EST	6441768
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Sequence 9822	AW177931	EST	6443968
Sequence 9823	AW178072	EST	6444109
Sequence 9824	AW194013	EST	6472742
Sequence 9825	AW237543	EST	6569932
Sequence 9826	AW239041	EST	6571431
Sequence 9827	AW239281	EST	6571671
Sequence 9828	AW245631	EST	6588624
Sequence 9829	AW245889	EST	6588882
Sequence 9830	AW245906	EST	6588899
Sequence 9831	AW245923	EST	6588916
	AW246049	EST	6589042
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Sequence 9833	AW246074		6589067
Sequence 9834	AW247459	EST	6590452
Sequence 9835	AW247494	EST	6590487
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Sequence 9837	AW250125	EST	6593118
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Sequence 9844	AW292897	EST	6699533
	AW293159	EST	6699795
Sequence 9845			
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Sequence 9847	AW294971	EST	6701607
Sequence 9848	AW296629	EST	6703265
Sequence 9849	AW303456	EST	6713145
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Sequence 9887	AW408423	EST	6927480
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Sequence 9889	AW410772	EST	6936313
Sequence 9890	AW411211	EST	6936752
Sequence 9891	AW411217	EST	6936758
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Sequence 9893	AW500117	EST	7112425
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Sequence 9896	AW501342	EST	7114784
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Sequence 9898	AW501390	EST	7114878
Sequence 9899	AW501401	EST	7114900
Sequence 9900	AW501421	EST	7114936
Sequence 9901	AW515416	EST	N/A
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Sequence 9969	W03691	EST	1275555
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Sequence 9971	W27890	EST	1307838
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Sequence 9975	W39482	EST	1321190
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Sequence 9990	Z78343		
Sequence 9991	Q46540	NUCPATENT	N/A
Sequence 9992	Q84560	NUCPATENT	N/A
Sequence 9993	T33624	NUCPATENT	N/A
Sequence 9994	T71086	NUCPATENT	N/A
Sequence 9995	T94119	NUCPATENT	N/A
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Sequence 9998	V34175	NUCPATENT	N/A
Sequence 9999	V34185	NUCPATENT	N/A
Sequence 10000	V35457	NUCPATENT	N/A
Sequence 10001	V36303	NUCPATENT	N/A
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Sequence 10003	V58631	NUCPATENT	N/A
Sequence 10004	V84411	NUCPATENT	N/A
Sequence 10005	V84546	NUCPATENT	N/A
Sequence 10006	V87058	NUCPATENT	N/A
Sequence 10007	V89685	NUCPATENT	N/A
Sequence 10008	V90259	NUCPATENT	N/A
Sequence 10009	V99828	NUCPATENT	N/A
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Sequence 10022	X98309	NUCPATENT	N/A
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Sequence 10029	Z15023	NUCPATENT	N/A
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Sequence 10033	Z38853	NUCPATENT	N/A
Sequence 10034	Z39903	NUCPATENT	N/A
Sequence 10035	Z41312	NUCPATENT	N/A
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Sequence 10040	Z60383	NUCPATENT	N/A
Sequence 10041	Z65271	NUCPATENT	N/A
Sequence 10042	Z80193	NUCPATENT	N/A
Sequence 10043	Z80472	NUCPATENT	N/A
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Sequence 10050	AC38120	PREPATNUC	N/A
Sequence 10051	AC38948	PREPATNUC	N/A
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Sequence 10047: Found in patent publication WO99/09061 CCCGCGGTGGCCGAGGTACTGCGTGGCCGTCCGCTGTTGTGTGGCCTGGGCCGCAA

Sequence 10048: Found in patent publication WO00/06698

CCGCGGTGGCGGCCGAGGTCTGAACTGCGGGGTCTCTATCGCACTGCTAGGGGTTCTGCT GCTGGGTGCGCGCGCCCCGCGCGGGGCAGAAGCTTTTGAGATTGCTCTGCCACGAGA AAGCAACATTACAGTTCTCATAAAGCTGGGGACCCCGACTCTGCTGGCAAAACCCTGTTA CATCGTCTTTTCTAAAAGACATATAACCATGTTGTCCATCAAGTCTGGAGAAAAGAATAGT CTTTACCTTTAGCTGCCAGAGTCCTGAGAATCACTTTGTCATANAGATCCAGAAAAATAT TGACTGTATGTCAGGCCCATGTCCTTTTGGGGAGGTTCAGCTTCAGCCCTCGACATCGTT GTTGCCTACCCTCAACAGAACTTTCATCTGGGATGTCAAA

Sequence 10049: Found in patent publication WO00/06698

TGATCTTCAGCTGCTCCTGAACTTTGGCACTGTAAGGGTTCTCTGTATCGTATGGTGGTG ACAAAGCTTTTAAAATAACCTNTCCTCCAGTGCATGGAAAAGGACTGGAGTATTTATNAA GT

Sequence 10050: Found in patent publication WO00/06730

ACTTGTGAGCCTGGACTGCCTTCCCCGGCCTCCTACATGAACCCCTTCCCGGTGCTCCAT
CTCATCGAGGACTTGAGGCTGGCCTTGGAGATGCTGGAGCTTCCTCAGGAGAGAGCAGCC
CTCCTGAGCCAGATCCCTGGCCCAACAGCTGCCTACATAAAGGAATGGTTTGAAGAGAGC
CTCCTGAGCCAGGTAAGCCACCACAGTGCTGCTAGTAATGAAACCTACCAGGAACGCTTGGCA
CGTCTAGAAGGGGATAAGGAGTCCCTCATATTGCAGGTGAGTGTCCTCACAGACCAAGTA
GAAGCCCAGGGAGAAAAGATTCGAGACCTGGAAGTGTCTTGGAAGGACCCAGGTGAAA
CTCAATGCTGCTGAAGAGATGCTTCAACAGGAGCTGCTAAGCCGCACATCTCTTTGAGACC
CAGAAGCTCGATCTGATGACTGAAGTGTCTGAGCTGAAGCTCAAGCTGGTTGGCATGGAG
AAGGAGCAGAGAGAGCAGAGAAAAGCAGAGAGAGTTACTGCAAGAGCTC
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Sequence 10051: Found in patent publication WO00/11171

Sequence 10052: Found in patent publication WO00/15799

ACCGCGGGGGCCGNGGNTTCNGCCTGGAGGAGCTCAGGGTGGCCGGCATTCACAAG

AGGTGGCCCGGACCATCGGCATTTCTGTGGATCCGAGGAGGCGGAACAAGTCCACGGAGTCCCTGCAGACCAACGTGCAGCGGCTGAAGGAGTACCACAAAGGGACCCAAATTCAGCGGTCTGTGCCTACAAACTTCATTAATAACTGCTTGCAGATTGGCAGCTATCTGGTCACTTGACATATCCAATGTTGCTATTTTGGTCTGGAGAAAGTTCTCCCTTTCTTCATCTACCTTAATTTCATGTCCATTTTTAAATAATTCAAACATTTTGGGGATGTCACGGCCAATGGAATTTCGAGAAAGCTTTGGAACAGCTACGTAAAAGTTCTCCTTAATCGATTAAGTTCAGGCATTATTTCTGGAACAGCTACGTAAAAGTCTGCAACAATTTCATCATCCCAAATCTTCTGTATCAGACTAAGTGCCTCCTGCAAATGCAGCTCCATTTTCTTCCGCTATTTTGACCTCTGATGCATTCTCTGTAAAAACACTAAAAACACACTGGTAAATGAGCAACTTAATTGATTTCCGAAGCCAATGGGTATGGCAAACTAAAAACACACTGGTAAATGGCTTCACGTTTTTCTTNTTTTCCAGTGCCATATNCAGTGTCAAAATCAA

ANAAAACACTTTGCTTTGGACTAGTTAAAGCCAANAATTTGA

Sequence 10053: Found in patent publication WO00/06698

TAGGGCGAATTGGAGCTCCCCGCGGTGGCGGCCGAGGTACCTGAGCCAGGAGGAGGCCC

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GGCCGTGACCAGGAGCTATTTAACGAATACCAGTTCAGCGTGGACCAACTTATGGAACT
GGCCGGGCTGAGCTGTGCTACAGCCATCGCCAAGGCATATCCCCCCACGTCCATGTCCAG
GAGCCCCCCTACTGTCCTGGTCATCTGTGGCCCGGGGAATAATGGAGGAGATGGTCTGGT
CTGTGCTCGACACCTCAAACTCTTTGGCTACGAGCCAACCATCTATTACCCCAAAAGGCC
TAACAAGCCCCTCTTCACTGCATTGGTGACCCAGTGTCAGAAAATGGACATCCCTTTCCT
TGGGGAAATGCCCGCAGAGCCCATGACGATTGATGAACTGTTGAGCTGGTGGATGCC
ATCTTTGGCTTCAGCTTCAAGGGCGATGTTCGGGAAC

Sequence 10047: Found in patent publication WO99/09061 CCCGCGGTGGCCGAGGTACTGCGTGGCCGTCCGCTGTTGTGTGGCCTGGGCCGCAA

Sequence 10048: Found in patent publication WO00/06698

Sequence 10049: Found in patent publication WO00/06698

TGATCTTCAGCTGCTCCTGAACTTTGGCACTGTAAGGGTTCTCTGTATCGTATGGTGGTG ACAAAGCTTTTAAAATAACCTNTCCTCCAGTGCATGGAAAAGGACTGGAGTATTTATNAA GT

Sequence 10050: Found in patent publication WO00/06730

Sequence 10051: Found in patent publication WO00/11171

Sequence 10052: Found in patent publication WO00/15799

ACCGCGGGGGCGGCCGNGGNTTCNGCCTGGAGGAGCTCAGGGTGGCCGGCATTCACAAG

AGGTGGCCCGGACCATCGGCATTTCTGTGGATCCGAGGAGGCGGAACAAGTCCACGGAGT
CCCTGCAGACCAACGTGCAGCGGCTGAAGGAGTACCACAAAGGGACCCAAATTCAGCGGT
CTGTGCCTACAAACTTCATTAATAACTGCTTGCAGATTGGCAGCTATCTGGTCACTTGAC
ATATCCAATGTTGCTATTTTGGTCTGGAGAAAGTTCTCCCTTTCTTCATCTACCTTAATT
TCATGTCCATTTTTAAATAATTCAAACATTTTGGGGATGTCACGGCCAATGGAATTTCGA
GAAAGCTTTGGATATTTTTTATTCAGTTTCTTCCTTAATCGATTAAGTTCAGGCATTATT
TCTGGAACAGCTACGTAAAAGTCTGCAACAATTTCATCATCCCAAATCTTCTGTATCAGA
CTAAGTGCCTCCTGCAAATGCAGCTCCATTTTCTTCCGCTATTTTGACCTCTGATGCATT
CTCTGTAAATACAGCAACTTTATTGATTTCCGAAGCCAATGGGTATGGCAAACTAANAAC
ACTGGTAAATGGCTTCACGTTTTTCTTNTTTTCCAGTGCCATATNCAGTGTCAAAATCAA

ANAAAACACTTTGCTTTGGACTAGTTAAAGCCAANAATTTGA

Sequence 10053: Found in patent publication WO00/06698

TAĞGGCGAATTGGAGCTCCCCGCGGTGGCGGCCGAGGTACCTGAGCCAGGAGGAGGCCC

GGCCGTGGACCAGGAGCTATTTAACGAATACCAGTTCAGCGTGGACCAACTTATGGAACT
GGCCGGGCTGAGCTGTGCTACAGCCATCGCCAAGGCATATCCCCCCACGTCCATGTCCAG
GAGCCCCCCTACTGTCCTGGTCATCTGTGGCCCGGGGAATAATGGAGGAGATGGTCTGGT
CTGTGCTCGACACCTCAAACTCTTTGGCTACGAGCCAACCATCTATTACCCCAAAAGGCC
TAACAAGCCCCTCTTCACTGCATTGGTGACCCAGTGTCAGAAAATGGACATCCCTTTCCT
TGGGGAAATGCCCGCAGAGCCCATGACGATTGATGAACTGTTGAGCTGGTGGTGGATGCC
ATCTTTGGCTTCAGCTTCAAGGGCGATGTTCGGGAAC

Sequence 10047: Found in patent publication WO99/09061 CCCGCGGTGGCCGAGGTACTGCGTGGCCGTCGCTGTTGTGTGGCCTGGCCGCAA

Sequence 10048: Found in patent publication WO00/06698

CCGCGGTGGCGGCCGAGGTCTGAACTGCGGGGTCTCTATCGCACTGCTAGGGGTTCTGCT GCTGGGTGCGCGCGCCCCGCGCGGGGCAGAAGCTTTTGAGATTGCTCTGCCACGAGA AAGCAACATTACAGTTCTCATAAAGCTGGGGACCCCGACTCTGCTGGCAAAACCCTGTTA CATCGTCTTTTCTAAAAGACATATAACCATGTTGTCCATCAAGTCTGGAGAAAAGAATAGT CTTTACCTTTAGCTGCCAGAGTCCTGAGAATCACTTTGTCATANAGATCCAGAAAAATAT TGACTGTATGTCAGGCCCATGTCCTTTTGGGGAGGTTCAGCTTCAGCCCTCGACATCGTT GTTGCCTACCCTCAACAGAACTTTCATCTGGGATGTCAAA

Sequence 10049: Found in patent publication WO00/06698

TGATCTTCAGCTGCTCCTGAACTTTGGCACTGTAAGGGTTCTCTGTATCGTATGGTGGTG ACAAAGCTTTTAAAATAACCTNTCCTCCAGTGCATGGAAAAGGACTGGAGTATTTATNAA

Sequence 10050: Found in patent publication WO00/06730

ACTTGTGAGCCTGGACTGGCTTCCCCGGCCTCCTACATGAACCCCTTCCCGGTGCTCCAT CTCATCGAGGACTTGAGGCTGGCCTTGGAGATGCTGGAGCTTCCTCAGGAGAGAGCAGCC CTCCTGAGCCAGATCCCTGGCCCAACAGCTGCCTACATAAAGGAATGGTTTGAAGAGAGC TTGTCCCAGGTAAGCCACCACAGTGCTGCTAGTAATGAAACCTACCAGGAACGCTTGGCA CGTCTAGAAGGGGATAAGGAGTCCCTCATATTGCAGGTGAGTGTCCTCACAGACCAAGTA GAAGCCCAGGGAGAAAAGATTCGAGACCTGGAAGTGTCTTGGAAGGACACCAGGTGAAA CTCAATGCTGCTGAAGAGATGCTTCAACAGGAGCTGCTAAGCCGCACATCTCTTGAGACC CAGAAGCTCGATCTGATGACTGAAGTGTCTGAGCTGAAGCTCAAGCTGGTTGGCATGGAG AAGGAGCAGAGAGAGCAGGAGAGAAAAGCAGAGAGATTACTGCAAGAGCTC AGGCCCCTCAAAATCAAAGTGGAAGAGTTTGGAAA

Sequence 10051: Found in patent publication WO00/11171

Sequence 10052: Found in patent publication WO00/15799

ACCGCGGGGGCCGNGGNTTCNGCCTGGAGGAGCTCAGGGTGGCCGGCATTCACAAG

AGGTGGCCCGGACCATCGGCATTTCTGTGGATCCGAGGAGGCGGAACAAGTCCACGGAGT CCCTGCAGACCAACGTGCAGCGGCTGAAGGAGTACCACAAAGGGACCCAAATTCAGCGGT CTGTGCCTACAAACTTCATTAATAACTGCTTGCAGATTGGCAGCTATCTGGTCACTTGAC ATATCCAATGTTGCTATTTTGGTCTGGAGAAAGTTCTCCCTTTCTTCATCTACCTTAATT TCATGTCCATTTTTAAATAATTCAAACATTTTGGGGATGTCACGGCCAATGGAATTTCGA GAAAGCTTTGGATATTTTTTATTCAGTTTCTTCCTTAATCGATTAAGTTCAGGCATTATT TCTGGAACAGCTACGTAAAAGTCTGCAACAATTTCATCATCCCAAATCTTCTGTATCAGA CTAAGTGCCTCCTGCAAATGCAGCTCCATTTTCTTCCGCTATTTTGACCTCTGATGCATT CTCTGTAAATACAGCAACTTTATTGATTTCCGAAGCCAATGGGTATGGCAAACTAANAAC ACTGGTAAATGGCTTCACGTTTTTCTTNTTTTCCAGTGCCATATNCAGTGTCAAAATCAA

ANAAAACACTTTGCTTTGGACTAGTTAAAGCCAANAATTTGA

Sequence 10053: Found in patent publication WO00/06698

ACATCTTCCAGGACAAGGTAAACTCTGACATGCACTAGGTATGTGCAGATCCCGGCCCCT GCCACCCAGCCTCATGCAAGTCATCCCCGACATGACCTTCACGACCGCAATGCAAGGAGG GGAAGAAGTCACAGCACTGATGAGGACAGCTGCAGAGGTGGCAGTGTGTGGACACAGGA AGTTTGGGCCCCTCCCTGCCCCAGCTTTCCTAGGCCAGAATTGTGTTTGGCAGTAATTG TCTGTTTAAAAAAATAAAAAGGAGAGGAAGCGTTCACCGCCACAAATCATAAAATGGACA TGACTGTGGAGTCTTACAGTTCAGGGTTCTTTCATTCACGTCCCTTCCTGTCTCGGTCTG CGGTCTTTACCACATCAATAGGACTTTTTATGCGTCCGGGTTAATTTTTCACTCCAGTGC GTCCTGTTGCAGGGACCGGAGCTGATGGGAGCTGCTTCTCCCCATGCCTCACTGGTCCCA GATCANGGCTTCAGGGACAGATGATGAGTCTCAAACGAGCCANCAGGGGTTCTTTTGGGT ATAAATGGGCAATTCGNCCTGTCTTAAGNCTGATGACCTCANCCGTGGTTTTTGGAT Sequence 10054: Found in patent publication WO00/18922

TAGGGCGAATTGGAGCTCCCCGCGGTGGCGGCCGAGGTACCTGAGCCAGGAGGAGGCCC

GGCCGTGGACCAGGAGCTATTTAACGAATACCAGTTCAGCGTGGACCAACTTATGGAACT GGCCGGGCTGAGCTGTGCTACAGCCATCGCCAAGGCATATCCCCCCACGTCCATGTCCAG GAGCCCCCTACTGTCCTGGTCATCTGTGGCCCGGGGAATAATGGAGGAGATGGTCTGGT CTGTGCTCGACACCTCAAACTCTTTGGCTACGAGCCAACCATCTATTACCCCAAAAGGCC TAACAAGCCCTCTTCACTGCATTGGTGACCCAGTGTCAGAAAATGGACATCCCTTTCCT TGGGGAAATGCCCGCAGAGCCCATGACGATTGATGAACTGTTGAGCTGGTGGTGGATGCC ATCTTTGGCTTCAGCTTCAAGGGCGATGTTCGGGAAC

Sequence 10047: Found in patent publication WO99/09061

CCCGCGGTGGCCGAGGTACTGCGTGGCCGTCCGCTGTTGTGTGGCCTGGGCCGCAA

Sequence 10048: Found in patent publication WO00/06698

Sequence 10049: Found in patent publication WO00/06698

TGATCTTCAGCTGCTCCTGAACTTTGGCACTGTAAGGGTTCTCTGTATCGTATGGTGGTG ACAAAGCTTTTAAAATAACCTNTCCTCCAGTGCATGGAAAAGGACTGGAGTATTTATNAA GT

Sequence 10050: Found in patent publication WO00/06730

ACTTGTGAGCCTGGACTGGCTTCCCCGGCCTCCTACATGAACCCCTTCCCGGTGCTCCAT CTCATCGAGGACTTGAGGCTTGCCCGGCCTCCTACATGAGCCCTCCTCAGGAGACTGCCTGGAGCTTGCTCAGGAGAGAGCAGCC CTCCTGAGCCAGATCCCTGGCCCAACAGCTGCCTACATAAAGGAATGGTTTGAAGAGAGC TTGTCCCAGGTAAGCCACCACAGTGCTGCTAGTAATGAAACCTACCAGGAACGCTTGGCA CGTCTAGAAGGGGATAAGGAGTCCCTCATATTGCAGGTGAGTGTCCTCACAGACCAAGTA GAAGCCCAGGGAAAAAGATTCGAGACCTGGAAGTGTCTTGGAAGGACACCAGGTGAAA CTCAATGCTGCTGAAGAGATGCTTCAACAGGAGCTGCTAAGCCGCACATCTCTTGAGACC CAGAAGCTCGATCTGATGACTGAAGTGTCTGAGCTGAAGCTCAAGCTGGTTGGCATGGAG AAGGAGCAGAGAGAGAAAAGCAGAGAGTTACTGCAAGAGCTC AGGCCCCTCAAAATCAAAGTGGAAGAGTTTGGAAA

Sequence 10051: Found in patent publication WO00/11171

Sequence 10052: Found in patent publication WO00/15799

ACCGCGGGGCCGNGGNTTCNGCCTGGAGGAGCTCAGGGTGGCCGGCATTCACAAG

AGGTGGCCCGGACCATCGGCATTTCTGTGGATCCGAGGAGGCGGAACAAGTCCACGGAGT
CCCTGCAGACCAACGTGCAGCGGCTGAAGGAGTACCACAAAGGGACCCAAATTCAGCGGT
CTGTGCCTACAAACTTCATTAATAACTGCTTGCAGATTGGCAGCTATCTGGTCACTTGAC
ATATCCAATGTTGCTATTTTGGTCTGGAGAAAGTTCTCCCTTTCTTCATCTACCTTAATT
TCATGTCCATTTTTAAATAATTCAAACATTTTGGGGATGTCACGGCCAATGGAATTTCGA
GAAAGCTTTGGATATTTTTATTCAGTTTCTTCCTTAATCGATTAAGTTCAGGCATTATT
TCTGGAACAGCTACGTAAAAGTCTGCAACAATTTCATCATCCCAAATCTTCTGTATCAGA
CTAAGTGCCTCCTGCAAATGCAGCTCCATTTTCTTCCGCTATTTTGACCTCTGATGCATT
CTCTGTAAATACAGCAACTTTATTGATTTCCGAAGCCAATGGGTATGGCAAACTAANAAC
ACTGGTAAATGGCTTCACGTTTTTCTTNTTTTCCAGTGCCATATNCAGTGTCAAAATCAA

ANAAAACACTTTGCTTTGGACTAGTTAAAGCCAANAATTTGA

Sequence 10053: Found in patent publication WO00/06698

ACATCTTCCAGGACAAGGTAAACTCTGACATGCACTAGGTATGTGCAGATCCCGGCCCCT
GCCACCCAGCCTCATGCAAGTCATCCCCGACATGACCTTCACGACCGCAATGCAAGGAGG
GGAAGAAAGTCACAGCACTGATGAGGACAGCTGCAGAGGTGGCAGTGTGTGGACACAGGA
AGTTTGGGCCCCCTGCCCCAGCTTTCCTAGGCCAGAATTGTGTTTGGCAGTAATTG
TCTGTTTAAAAAAATAAAAAGGAGAGGAAGCGTTCACCGCCACAAATCATAAAATGGACA
TGACTGTGGAGTCTTACAGTTCAGGGTTCTTTCATTCACGTCCCTTCCTGTCTCGGTCTG
CGGTCTTTACCACATCAATAGGACTTTTTATGCGTCCGGGTTAATTTTTCACTCCAGTGC
GTCCTGTTGCAGGGACCGGAGCTGATGGGAGCTGCTTCTCCCCATGCCTCACTGGTCCCA
GATCANGGCTTCAGGGACAGATGATGAGTCTCAAACGAGCCANCAGGGGTTCTTTTGGGT
ATAAATGGGCAATTCGNCCTGTCTTAAGNCTGATGACCTCANCCGTGGTTTTTTGGAT
Sequence 10054: Found in patent publication WO00/18922

TAGGGCGAATTGGAGCTCCCCGCGGTGGCGGCCGAGGTACCTGAGCCAGGAGGAGGCCC

GGCCGTGACCAGGAGCTATTTAACGAATACCAGTTCAGCGTGGACCAACTTATGGAACT
GGCCGGGCTGAGCTGTGCTACAGCCATCGCCAAGGCATATCCCCCCACGTCCATGTCCAG
GAGCCCCCCTACTGTCCTGGTCATCTGTGGCCCGGGGAATAATGGAGGAGATGGTCTGGT
CTGTGCTCGACACCTCAAACTCTTTGGCTACGAGCCAACCATCTATTACCCCAAAAGGCC
TAACAAGCCCCTCTTCACTGCATTGGTGACCCAGTGTCAGAAAATGGACATCCCTTTCCT
TGGGGAAATGCCCGCAGAGCCCATGACGATTGATGAACTGTTGAGCTGGTGGTGATGCC
ATCTTTGGCTTCAGCTTCAAGGGCGATGTTCGGGAAC

Sequence 10047: Found in patent publication WO99/09061

CCCGCGGTGGCCGAGGTACTGCGTGGCCGTCCGCTGTTGTGTGGCCTGGGCCGCAA

Sequence 10048: Found in patent publication WO00/06698

Sequence 10049: Found in patent publication WO00/06698

TGATCTTCAGCTGCTCCTGAACTTTGGCACTGTAAGGGTTCTCTGTATCGTATGGTGGTG ACAAAGCTTTTAAAATAACCTNTCCTCCAGTGCATGGAAAAGGACTGGAGTATTTATNAA GT

Sequence 10050: Found in patent publication WO00/06730

Sequence 10051: Found in patent publication WO00/11171

Sequence 10052: Found in patent publication WO00/15799

ACCGCGGGGGCCGNGGNTTCNGCCTGGAGGAGCTCAGGGTGGCCGGCATTCACAAG

ANAAAACACTTTGCTTTGGACTAGTTAAAGCCAANAATTTGA

Sequence 10053: Found in patent publication WO00/06698

TAĠGGCGAATTGGAGCTĊCCCĠCGGTGGCGGCCGAGGTACCTGAGCCAGGAGGAGGCCC

GGCCGTGGACCAGGAGCTATTTAACGAATACCAGTTCAGCGTGGACCAACTTATGGAACT
GGCCGGGCTGAGCTGTCCTACAGCCATCGCCAAGGCATATCCCCCCACGTCCATGTCCAG
GAGCCCCCCTACTGTCCTGGTCATCTGTGGCCCGGGGAATAATGGAGGAGATGGTCTGGT
CTGTGCTCGACACCTCAAACTCTTTGGCTACGAGCCAACCATCTATTACCCCAAAAGGCC
TAACAAGCCCCTCTTCACTGCATTGGTGACCCAGTGTCAGAAAATGGACATCCCTTTCCT
TGGGGAAATGCCCGCAGAGCCCATGACGATTGATGAACTGTTGAGCTGGTGGTGGATGCC
ATCTTTGGCTTCAGCTTCAAGGGCGATGTTCGGGAAAC

Sequence 10047: Found in patent publication WO99/09061

CCCGCGTGGCCGAGGTACTGCGTGGCCGTCCGCTGTTGTGTGGCCTGGGCCGCAA

Α

Sequence 10048: Found in patent publication WO00/06698

CCGCGGTGGCGGCCGAGGTCTGAACTGCGGGGTCTCTATCGCACTGCTAGGGGTTCTGCT GCTGGGTGCGCGCGCCCGCGCGCGGGGCAGAAGCTTTTGAGATTGCTCTGCCACGAGA AAGCAACATTACAGTTCTCATAAAGCTGGGGACCCCGACTCTGCTGGCAAAACCCTGTTA CATCGTCTTTTCTAAAAGACATATAACCATGTTGTCCATCAAGTCTGGAGAAAAGAATAGT CTTTACCTTTAGCTGCCAGAGTCCTGAGAATCACTTTGTCATANAGATCCAGAAAAATAT TGACTGTATGTCAGGCCCATGTCCTTTTGGGGAGGTTCAGCTTCAGCCCTCGACATCGTT GTTGCCTACCCTCAACAGAACTTTCATCTGGGATGTCAAA

Sequence 10049: Found in patent publication WO00/06698

TGATCTTCAGCTGCTCCTGAACTTTGGCACTGTAAGGGTTCTCTGTATCGTATGGTGGTG ACAAAGCTTTTAAAATAACCTNTCCTCCAGTGCATGGAAAAGGACTGGAGTATTTATNAA GT

Sequence 10050: Found in patent publication WO00/06730

Sequence 10051: Found in patent publication WO00/11171

Sequence 10052: Found in patent publication WO00/15799

ACCGCGGGGCCGNGGNTTCNGCCTGGAGGAGCTCAGGGTGGCCGGCATTCACAAG

Α

AGGTGGCCCGGACCATCGGCATTTCTGTGGATCCGAGGAGGCGGAACAAGTCCACGGAGT CCCTGCAGACCAACGTGCAGCGGCTGAAGGAGTACCACAAAGGGACCCAAATTCAGCGGT CTGTGCCTACAAACTTCATTAATAACTGCTTGCAGATTGGCAGCTATCTGGTCACTTGAC ATATCCAATGTTGCTATTTTGGTCTGGAGAAAGTTCTCCCTTTCTTCATCTACCTTAATT TCATGTCCATTTTTAAATAATTCAAACATTTTGGGGATGTCACGGCCAATGGAATTTCGA GAAAGCTTTGGATATTTTTTATTCAGTTTCTTCCTTAATCGATTAAGTTCAGGCATTATT TCTGGAACAGCTACGTAAAAGTCTGCAACAATTTCATCATCCCAAATCTTCTGTATCAGA CTAAGTGCCTCCTGCAAATGCAGCTCCATTTTCTTCCGCTATTTTGACCTCTGATGCATT CTCTGTAAATACAGCAACTTTATTGATTTCCGAAGCCAATGGGTATGGCAAACTAANAAC ACTGGTAAATGGCTTCACGTTTTTCTTNTTTTCCAGTGCCATATNCAGTGTCAAAATCAA

ANAAAACACTTTGCTTTGGACTAGTTAAAGCCAANAATTTGA
Sequence 10053: Found in patent publication WO00/06698
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GATCANGGCTTCAGGGACAGATGATGAGTCTCAAACGAGCCANCAGGGGTTCTTTTGGGT
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Sequence 10054: Found in patent publication WO00/18922

TAGGGCGAATTGGAGCTCCCCGCGGTGGCGGCCGAGGTACCTGAGCCAGGAGGAGGCCC

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GAGCCCCCTACTGTCCTGGTCATCTGTGGCCCGGGGAATAATGGAGGAGATGGTCTGGT
CTGTGCTCGACACCTCAAACTCTTTGGCTACGAGCCAACCATCTATTACCCCAAAAGGCC
TAACAAGCCCCTCTTCACTGCATTGGTGACCCAGTGTCAGAAAATGGACATCCCTTTCCT
TGGGGAAATGCCCGCAGAGCCCATGACGATTGATGAACTGTTGAGCTGGTGGTGGATGCC
ATCTTTGGCTTCAGCTTCAAGGGCGATGTTCGGGAAC

Sequence 10047: Found in patent publication WO99/09061 CCCGCGGTGGCCGAGGTACTGCGTGGCCGTCGCTGTTGTGTGGCCTGGCCGCAA

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Sequence 10048: Found in patent publication WO00/06698

Sequence 10049: Found in patent publication WO00/06698

TGATCTTCAGCTGCTCCTGAACTTTGGCACTGTAAGGGTTCTCTGTATCGTATGGTGGTG ACAAAGCTTTTAAAATAACCTNTCCTCCAGTGCATGGAAAAGGACTGGAGTATTTATNAA GT

Sequence 10050: Found in patent publication WO00/06730

Sequence 10051: Found in patent publication WO00/11171

Sequence 10052: Found in patent publication WO00/15799

ACCGCGGGGGCCGNGGNTTCNGCCTGGAGGAGCTCAGGGTGGCCGGCATTCACAAG

ANAAAACACTTTGCTTTGGACTAGTTAAAGCCAANAATTTGA

Sequence 10053: Found in patent publication WO00/06698

ATCTTTGGCTTCAGCTTCAAGGGCGATGTTCGGGAAC

TAGGGCGAATTGGAGCTCCCCGCGGTGGCGGCCGAGGTACCTGAGCCAGGAGGAGGCCC

GGCCGTGGACCAGGAGCTATTTAACGAATACCAGTTCAGCGTGGACCAACTTATGGAACT
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Sequence 10047: Found in patent publication WO99/09061

CCCGCGGTGGCCGAGGTACTGCGTGGCCGTCCGCTGTTGTGTGGCCTGGGCCGCAA

Α

Sequence 10048: Found in patent publication WO00/06698

Sequence 10049: Found in patent publication WO00/06698

TGATCTTCAGCTGCTCCTGAACTTTGGCACTGTAAGGGTTCTCTGTATCGTATGGTGGTG ACAAAGCTTTTAAAATAACCTNTCCTCCAGTGCATGGAAAAGGACTGGAGTATTTATNAA GT

Sequence 10050: Found in patent publication WO00/06730

ACTTGTGAGCCTGGACTGGCTTCCCGGCCTCCTACATGAACCCCTTCCCGGTGCTCCAT CTCATCGAGGACTTGAGGCTGGCCTTGGAGATGCTGGAGCTTCCTCAGGAGAGAGCAGCC CTCCTGAGCCAGATCCCTGGCCCAACAGCTGCCTACATAAAGGAATGGTTTGAAGAGAGC TTGTCCCAGGTAAGCCACCACAGTGCTGCTAGTAATGAAACCTACCAGGAACGCTTGGCA CGTCTAGAAGGGGATAAGGAGTCCCTCATATTGCAGGTGAGTGTCCTCACAGACCAAGTA GAAGCCCAGGGAGAAAAGATTCGAGACCTGGAAGTGTCTTGGAAGGACACCAGGTGAAA CTCAATGCTGCTGAAGAGATGCTTCAACAGGAGCTGCTAAGCCGCACATCTCTTGAGACC CAGAAGCTCGATCTGATGACTGAAGTGTCTGAGCTGAAGCTCAAGCTGGTTGGCATGGAG AAGGAGCAGAGAGAGAGAGAGAGAGAGAAAAGCAGAGAGAGAGAGAGAGCTCAAGAGCTCAAAATCAAAGTGGAAGAGTTTGGAAA

Sequence 10051: Found in patent publication WO00/11171

Sequence 10052: Found in patent publication WO00/15799

ACCGCGGGGGCGGCCGNGGNTTCNGCCTGGAGGAGCTCAGGGTGGCCGGCATTCACAAG

AGGTGGCCCGGACCATCGGCATTTCTGTGGATCCGAGGAGGCGGAACAAGTCCACGGAGT
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TCATGTCCATTTTTAAATAATTCAAACATTTTGGGGATGTCACGGCCAATGGAATTTCGA
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CTCTGTAAATACAGCAACTTTATTGATTTCCGAAGCCAATGGGTATGGCAAACTAANAAC
ACTGGTAAATGGCTTCACGTTTTTCTTNTTTTCCAGTGCCATATNCAGTGTCAAAATCAA

ANAAAACACTTTGCTTTGGACTAGTTAAAGCCAANAATTTGA

ATAAATGGGCAATTCGNCCTGTCTTAAGNCTGATGACCTCANCCGTGGTTTTTGGAT Sequence 10054: Found in patent publication WO00/18922 TAGGGCGAATTGGAGCTCCCCGCGGTGGCGGCCGAGGTACCTGAGCCAGGAGGAGGCCC

GATCANGGCTTCAGGGACAGATGATGAGTCTCAAACGAGCCANCAGGGGTTCTTTTGGGT

GGCCGTGACCAGGAGCTATTTAACGAATACCAGTTCAGCGTGGACCAACTTATGGAACT
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CTGTGCTCGACACCTCAAACTCTTTGGCTACGAGCCAACCATCTATTACCCCAAAAGGCC
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<u>Table 3-5</u>

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•		Genbank	4156135
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Sequence 10309	AC007684	Genbank	5836173
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Sequence 11365	AA935796	EST	
Sequence 11366	AA939273	EST	3092953
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Sequence 11369	AA962465	EST	3134629
Sequence 11370	AA973265	EST	3148445
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Sequence 11372	AA977036	EST	3154482
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Sequence 11391	Al039136	EST	3278330
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Sequence 11393	Al040803	EST	3278925 3279997
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Sequence 11395	Al051985 Al052598	EST	3307976 3308589
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Sequence 11467	Al284545	EST	3922778
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Sequence 11538	A1858089	EST	5511705
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Sequence 11561	AL134867	EST	6603054
Sequence 11562	AL135047	EST	6603234
Sequence 11563	AL138273	EST	6854954
Sequence 11564	AL138451	EST	6855132
Sequence 11565	AW003648	EST	5850564
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Sequence 11566	AW008029	EST	5856807
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Sequence 11568	AW008782	EST	5857560
Sequence 11569	AW021322	EST	5874782
Sequence 11570	AW021588	EST	5875118
Sequence 11571	AW022159	EST	5875689
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Sequence 11574	AW069285	EST	6024283
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Sequence 11576	AW073580	EST	6028578
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Sequence 11583	AW157858	EST	6229259
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Sequence 11585	AW161328	EST	6300361
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Sequence 11594	AW247236	EST	6590229
Sequence 11595	AW250104	EST	
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Sequence 11597	AW294626		6701242
•	AW297943	EST	6701262
Sequence 11598		EST	6704579
Sequence 11599	AW302841	EST	6712521
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Sequence 11602	AW363975	EST	6868625
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Sequence 11604	AW365960	EST	6870610
Sequence 11605	AW368992	EST	6873642
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Sequence 11616	AW503119	EST	7118224
Sequence 11617	AW518064	EST	N/A
Sequence 11618	AW573593	EST	7238326
Sequence 11619	AW574740	EST	7246279
Sequence 11620	AW575094	EST	7246633
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<u>Table 3-5</u>

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Sequence 11627	F20959	EST	2060135
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Sequence 11632	H16546	EST	882771
Sequence 11633	H18560	EST	884800
Sequence 11634	H22094	EST	890789
Sequence 11635	H40387	EST	916439
Sequence 11636	H54240	EST	994387
Sequence 11637	H68450	EST	1027190
Sequence 11638	H90819	EST	1081249
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Sequence 11642	N21624	EST	1126794
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Sequence 11674	Z33656	NUCPATENT	488829
Sequence 11675	Z39728	NUCPATENT	565523
Sequence 11676	AC02994	PREPATNUC	N/A
Sequence 11677	AC31122	PREPATNUC	N/A
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Sequence 11679	AC35230	PREPATNUC	N/A
Sequence 11680	AC36946	PREPATNUC	N/A
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Sequence 11682	A16794	Genbank	512417
Sequence 11683	A17003	Genbank	512419
Sequence 11684	A20255	Genbank	1247401
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Sequence 11687	AB001535	Genbank	3928755
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Sequence 11694	AB003102	Genbank	1945608
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Sequence		AC005042	Genbank	4156138
Sequence		AC005084	Genbank	3659503
Sequence		AC005089	Genbank	5732140
Sequence		AC005009 AC005104	Genbank	4218027
Sequence		AC005104 AC005153	Genbank	3242766
•			Genbank	
Sequence		AC005210		6249673
Sequence		AC005257	Genbank	3289994
Sequence		AC005263	Genbank	3289978
Sequence		AC005318	Genbank	3885345
Sequence		AC005324	Genbank	3366582
Sequence		AC005329	Genbank	3342732
Sequence		AC005332	Genbank	3659494
Sequence		AC005369	Genbank	3367505
Sequence		AC005382	Genbank	3386587
Sequence	11865	AC005488	Genbank	5836194
Sequence	11866	AC005550	Genbank	3478668
Sequence	11867	AC005614	Genbank	3540175
Sequence	11868	AC005621	Genbank	3548788
Sequence	11869	AC005627	Genbank	7243867
Sequence	11870	AC005629	Genbank	7243877
Sequence	11871	AC005630	Genbank	4159882
Sequence	11872	AC005661	Genbank	3818348
Sequence	11873	AC005702	Genbank	3688100
Sequence		AC005726	Genbank	3810672
Sequence		AC005786	Genbank	3702287
Sequence		AC005789	Genbank	3702281
Sequence		AC005837	Genbank	3849820
Sequence		AC005838	Genbank	3947427
Sequence		AC005841	Genbank	4731044
Sequence		AC005912	Genbank	4165005
Sequence		AC005971	Genbank	4314421
Sequence		AC006008	Genbank	5091655
Sequence		AC006014	Genbank	4699959
Sequence		AC006017	Genbank	4508141
Sequence		AC006040	Genbank	4314426
Sequence		AC006057	Genbank	4731048
Sequence :		AC006115	Genbank	3962498
Sequence '		AC006137	Genbank	9887710
Sequence :		AC006160	Genbank	
Sequence :		AC006100 AC006213	_	5701616
Sequence :		AC006213 AC006230	Genbank Genbank	4160143
Sequence :		AC006255		4966389
		AC006253 AC006273	Genbank	5306223
Sequence :			Genbank	4096060
Sequence :		AC006299	Genbank	4106997
Sequence :		AC006475	Genbank	4753283
Sequence :		AC006477	Genbank	7243871
Sequence 1		AC006480	Genbank	6289252
Sequence 1		AC006512	Genbank	4926863
Sequence 1		AC006566	Genbank	4309780
Sequence 1		AC006581	Genbank	4914350
Sequence 1	11901	AC007032	Genbank	5523832

Sequence 11902	AC007041	Genbank	5708471
	AC007055	Genbank	4885691
Sequence 11903			
Sequence 11904	AC007066	Genbank	4508098
Sequence 11905	AC007073	Genbank	4416547
Sequence 11906	AC007172	Genbank	4731066
Sequence 11907	AC007191	Genbank	4558643
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Sequence 11909	AC007253	Genbank	5649375
Sequence 11910	AC007384	Genbank	5708473
Sequence 11911	AC007406	Genbank	4689442
Sequence 11912	AC007528	Genbank	5263309
Sequence 11913	AC007537	Genbank	4914348
Sequence 11914	AC007541	Genbank	4982536
Sequence 11915	AC007564	Genbank	4982534
Sequence 11916	AC007565	Genbank	4835751
Sequence 11917	AC007736	Genbank	5732177
_			5306288
Sequence 11918	AC007938	Genbank	
Sequence 11919	AC007999	Genbank	7007656
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Sequence 11921	AC008101	Genbank	6970743
Sequence 11922	AC008102	Genbank	9797815
Sequence 11923	AC008122	Genbank	5931373
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Sequence 11926	AC008498	Genbank	6850299
Sequence 11927	AC008893	Genbank	7259691
Sequence 11928	AC009330	Genbank	6553960
Sequence 11929	AC010206	Genbank	6539154
Sequence 11930	AC010582	Genbank	6721135
	AC011331	Genbank	9929686
Sequence 11931			
Sequence 11932	AC012087	Genbank	7114465
Sequence 11933	AC020663	Genbank	6682593
Sequence 11934	AC021049	Genbank	7248920
Sequence 11935	AF001549	Genbank	3355302
Sequence 11936	AF001977	Genbank	2914738
Sequence 11937	AF002668	Genbank	2232173
	AF003837	Genbank	2228792
Sequence 11938			
Sequence 11939	AF004162	Genbank	3046385
Sequence 11940	AF004339	Genbank	2245559
Sequence 11941	AF004341	Genbank	2245563
Sequence 11942	AF004342	Genbank	2245565
Sequence 11943	AF005896	Genbank	2209244
Sequence 11944	AF006083	Genbank	2282031
		Genbank	
Sequence 11945	AF007544		2970122
Sequence 11946	AF007835	Genbank	4102034
Sequence 11947	AF008303	Genbank	2580619
Sequence 11948	AF011889	Genbank	2335186
Sequence 11949	AF012023	Genbank	2305237
Sequence 11950	AF012072	Genbank	9967556
		Genbank	2832227
Sequence 11951	AF012126		
Sequence 11952	AF015812	Genbank	2599359
Sequence 11953	AF016709	Genbank	2731560
Sequence 11954	AF019413	Genbank	2347130
Sequence 11955	AF020057	Genbank	3150436
Sequence 11956	AF020351	Genbank	2655052
Sequence 11957	AF020352	Genbank	2655054
2242200 . 1001	0_000_		

Sequence 11958	AF021819	Genbank	2460317
Sequence 11959	AF022913	Genbank	2558890
		Genbank	2559009
Sequence 11960	AF026292		
Sequence 11961	AF026816	Genbank	2583078
Sequence 11962	AF028832	Genbank	3287488
Sequence 11963	AF030186	Genbank	3831546
Sequence 11964	AF032387	Genbank	2641556
		Genbank	2645728
Sequence 11965	AF033095		
Sequence 11966	AF035840	Genbank	3800739
Sequence 11967	AF036715	Genbank	4007385
Sequence 11968	AF038178	Genbank	2795898
Sequence 11969	AF038200	Genbank	2795921
	AF038554	Genbank	2895558
Sequence 11970			3252876
Sequence 11971	AF038633	Genbank	
Sequence 11972	AF038955	Genbank	3329379
Sequence 11973	AF041483	Genbank	3493528
Sequence 11974	AF042166	Genbank	3298596
Sequence 11975	AF042836	Genbank	3776241
Sequence 11976	AF044957	Genbank	4164445
•			4164447
Sequence 11977	AF044958	Genbank	
Sequence 11978	AF047185	Genbank	2909861
Sequence 11979	AF047440	Genbank	3335135
Sequence 11980	AF047472	Genbank	2921872
Sequence 11981	AF049105	Genbank	2984656
Sequence 11982	AF051316	Genbank	6580759
		Genbank	3095110
Sequence 11983	AF051894		
Sequence 11984	AF052129	Genbank	3360438
Sequence 11985	AF052642	Genbank	2981453
Sequence 11986	AF053356	Genbank	3135305
Sequence 11987	AF054984	Genbank	3005692
Sequence 11988	AF054990	Genbank	3005703
Sequence 11989	AF055473	Genbank	nil
		Genbank	nil
Sequence 11990	AF055474		
Sequence 11991	AF061737	Genbank	4335938
Sequence 11992	AF064019	Genbank	3347856
Sequence 11993	AF064084	Genbank	3135668
Sequence 11994	AF064245	Genbank	3930528
Sequence 11995	AF064861	Genbank	3171154
	AF067517	Genbank	3850313
Sequence 11996		Gondann	
Sequence 11997	AF067844	Genbank	4240386
Sequence 11998	AF068754	Genbank	3283408
Sequence 11999	AF069378	Genbank	3982733
Sequence 12000	AF070646	Genbank	3283920
Sequence 12001	AF070649	Genbank	3283923
Sequence 12002	AF070652	Genbank	4454679
		Genbank	4454685
Sequence 12003	AF070655		
Sequence 12004	AF073792	Genbank	3300101
Sequence 12005	AF075053	Genbank	3377594
Sequence 12006	AF075242	Genbank	4377992
Sequence 12007	AF077042	Genbank	4689131
Sequence 12008	AF077043	Genbank	4689133
Sequence 12009	AF077046	Genbank	4689139
		Genbank	4689151
Sequence 12010	AF077052		
Sequence 12011	AF077202	Genbank	4679017
Sequence 12012	AF078855	Genbank	5531824
Sequence 12013	AF081282	Genbank	4336324
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<u>Table 3-5</u>

Sequence 12014	AF083441	Genbank	5813822
Sequence 12015	AF085361	Genbank	5114056
Sequence 12016	AF086168	Genbank	3483513
Sequence 12017	AF086169	Genbank	3483514
Sequence 12018	AF086210	Genbank	3483555
Sequence 12019	AF086236	Genbank	3483581
Sequence 12020	AF086329	Genbank	3483674
Sequence 12021	AF086465	Genbank	3483810
Sequence 12022	AF086557	Genbank	3483902
	AF091075	Genbank	3859987
Sequence 12023	AF091073	Genbank	3860003
Sequence 12024			3860005
Sequence 12025	AF091084	Genbank	
Sequence 12026	AF092124	Genbank	3659900
Sequence 12027	AF092135	Genbank	5138919
Sequence 12028	AF098162	Genbank	3929582
Sequence 12029	AF098865	Genbank	4204674
Sequence 12030	AF100748	Genbank	5410281
Sequence 12031	AF100928	Genbank	4323586
Sequence 12032	AF103801	Genbank	6048969
Sequence 12033	AF105278	Genbank	4324471
Sequence 12034	AF107893	Genbank	4008128
Sequence 12035	AF109872	Genbank	4028621
Sequence 12036	AF110304	Genbank	7106722
Sequence 12037	AF110460	Genbank	4324698
Sequence 12038	AF110774	Genbank	6523794
Sequence 12039	AF111847	Genbank	7211441
Sequence 12040	AF112215	Genbank	6563217
Sequence 12041	AF112222	Genbank	6563229
Sequence 12042	AF113016	Genbank	6642755
Sequence 12043	AF113680	Genbank	6855605
Sequence 12044	AF113700	Genbank	6855634
Sequence 12045	AF117235	Genbank	6563243
Sequence 12046	AF117829	Genbank	4151947
Sequence 12047	AF118091	Genbank	6650827
	AF119663	Genbank	6563251
Sequence 12048	AF121219	Genbank	4903275
Sequence 12049		Genbank	
Sequence 12050	AF122004	G. G. 1 1 2 1 1 1 1 1	5059061
Sequence 12051	AF128536	Genbank	5305705
Sequence 12052	AF129927	Genbank	4558506
Sequence 12053	AF130249	Genbank	4455134
Sequence 12054	AF131738	Genbank	4406548
Sequence 12055	AF131838	Genbank	4406677
Sequence 12056	AF131856	Genbank	4406702
Sequence 12057	AF132965	Genbank	4680700
Sequence 12058	AF132973	Genbank	4680716
Sequence 12059	AF139065	Genbank	6635330
Sequence 12060	AF144700	Genbank	5107075
Sequence 12061	AF144755	Genbank	5006628
Sequence 12062	AF146191	Genbank	5678818
Sequence 12063	AF147330	Genbank	4761681
Sequence 12064	AF150089	Genbank	5107165
Sequence 12065	AF151020	Genbank	7106761
Sequence 12066	AF151066	Genbank	7106853
Sequence 12067	AF151109	Genbank	6166337
Sequence 12068	AF151807	Genbank	4929566
Sequence 12069	AF151832	Genbank	4929616
Coquence 12000	/11 10 100Z	Gonbank	.5200.0

Sequence 12070	AF151841	Genbank	4929634
Sequence 12071	AF151861	Genbank	4929674
Sequence 12072	AF151875	Genbank	4929702
			4929710
Sequence 12073	AF151879	Genbank	
Sequence 12074	AF151884	Genbank	4929720
Sequence 12075	AF151886	Genbank	4929724
Sequence 12076	AF151889	Genbank	4929730
Sequence 12077	AF151893	Genbank	4929738
•		Genbank	4929762
Sequence 12078	AF151905		
Sequence 12079	AF151906	Genbank	4929764
Sequence 12080	AF152462	Genbank	5565976
Sequence 12081	AF155568	Genbank	5031511
Sequence 12082	AF156965	Genbank	5731112
Sequence 12083	AF161389	Genbank	6841191
Sequence 12084	AF161448	Genbank	6841309
	AF161466	Genbank	6841455
Sequence 12085			
Sequence 12086	AF161500	Genbank	6841523
Sequence 12087	AF161503	Genbank	6841529
Sequence 12088	AF161512	Genbank	6841547
Sequence 12089	AF161556	Genbank	6841379
Sequence 12090	AF165926	Genbank	5817858
Sequence 12091	AF168787	Genbank	7239175
	AF168956	Genbank	5702387
Sequence 12092		Genbank	7108914
Sequence 12093	AF174496	.	
Sequence 12094	AF176574	Genbank	5762481
Sequence 12095	AF179274	Genbank	5911402
Sequence 12096	AF187320	Genbank	6164847
Sequence 12097	AF187554	Genbank	6653225
Sequence 12098	AF188178	Genbank	6164864
Sequence 12099	AF188745	Genbank	6425043
Sequence 12100	AF191018	Genbank	6457339
	AF191298	Genbank	7656642
Sequence 12101			
Sequence 12102	AF196969	Genbank	6289073
Sequence 12103	AF201077	Genbank	6456748
Sequence 12104	AF203815	Genbank	6979641
Sequence 12105	AF205588	Genbank	6531675
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Sequence 12107	AF207550	Genbank	6470333
Sequence 12108	AF217403	Genbank	6691160
Sequence 12109	AF224669	Genbank	7012905
	AJ001981	Genbank	2546978
Sequence 12110			
Sequence 12111	AJ010069	Genbank	3483012
Sequence 12112	AJ010341	Genbank	4490794
Sequence 12113	AJ011803	Genbank	4753760
Sequence 12114	AJ012499	Genbank	5441359
Sequence 12115	AJ131693	Genbank	4584422
Sequence 12116	AJ224442	Genbank	2911586
Sequence 12117	AJ243669	Genbank	6688156
	AJ249731	Genbank	5921469
Sequence 12118			
Sequence 12119	AJ250042	Genbank	6013005
Sequence 12120	AJ250915	Genbank	6996445
Sequence 12121	AK000049	Genbank	7019880
Sequence 12122	AK000052	Genbank	7019885
Sequence 12123	AK000068	Genbank	7019914
Sequence 12124	AK000083	Genbank	7019940
Sequence 12125	AK000171	Genbank	7020084
5544550 IE IEO			

Sequence 12126	AK000221	Genbank	7020163
Sequence 12127	AK000303	Genbank	7020295
		Genbank	7020324
Sequence 12128	AK000319		
Sequence 12129	AK000395	Genbank	7020455
Sequence 12130	AK000400	Genbank	7020463
Sequence 12131	AK000434	Genbank	7020520
Sequence 12132	AK000462	Genbank	7020566
Sequence 12133	AK000470	Genbank	7020580
Sequence 12134	AK000526	Genbank	7020680
Sequence 12135	AK000560	Genbank	7020738
Sequence 12136	AK000615	Genbank	7020827
Sequence 12137	AK000620	Genbank	7020834
		Genbank	7020839
Sequence 12138	AK000624		
Sequence 12139	AK000656	Genbank	7020889
Sequence 12140	AK000667	Genbank	7020906
Sequence 12141	AK000669	Genbank	7020909
Sequence 12142	AK000818	Genbank	7021128
Sequence 12143	AK000931	Genbank	7021906
Sequence 12144	AK001067	Genbank	7022104
Sequence 12145	AK001088	Genbank	7022136
Sequence 12146	AK001163	Genbank	7022244
Sequence 12147	AK001192	Genbank	7022292
Sequence 12148	AK001221	Genbank	7022340
Sequence 12149	AK001321	Genbank	7022503
•	AK001321	Genbank	7022533
Sequence 12150			
Sequence 12151	AK001354	Genbank	7022561
Sequence 12152	AK001408	Genbank	7022645
Sequence 12153	AK001441	Genbank	7022699
Sequence 12154	AK001453	Genbank	7022720
Sequence 12155	AK001489	Genbank	7022777
Sequence 12156	AK001510	Genbank	7022808
	AK001516	Genbank	7022851
Sequence 12157			
Sequence 12158	AK001564	Genbank	7022893
Sequence 12159	AK001614	Genbank	7022977
Sequence 12160	AK001689	Genbank	7023104
Sequence 12161	AK001718	Genbank	7023153
Sequence 12162	AK001764	Genbank	7023235
Sequence 12163	AK001780	Genbank	7023265
	AK001700 AK001902	Genbank	7023455
Sequence 12164			
Sequence 12165	AK002031	Genbank	7023667
Sequence 12166	AK002100	Genbank	7023777
Sequence 12167	AL008721	Genbank	3171883
Sequence 12168	AL008723	Genbank	5101762
Sequence 12169	AL009179	Genbank	3217024
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Sequence 12171	AL020997	Genbank	2804155
Sequence 12172	AL021368	Genbank	3080468
Sequence 12173	AL021397	Genbank	3319943
Sequence 12174	AL021707	Genbank	4582132
Sequence 12175	AL021917	Genbank	9367203
Sequence 12176	AL022153	Genbank	3319674
Sequence 12177	AL022238	Genbank	4176442
Sequence 12178	AL022240	Genbank	4826471
Sequence 12179	AL022313	Genbank	4200326
Sequence 12180	AL022326	Genbank	3550039
Sequence 12181	AL022718	Genbank	3763969
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Sequence 12182	AL022725	Genbank	5679748
Sequence 12183	AL023581	Genbank	3445283
Sequence 12184	AL023582	Genbank	3805923
Sequence 12185	AL023803	Genbank	9408726
Sequence 12186	AL023807	Genbank	5679750
Sequence 12187	AL024474	Genbank	3395511
Sequence 12188	AL031005	Genbank	3287156
Sequence 12189	AL031256	Genbank	3738133
Sequence 12190	AL031274	Genbank	3421083
Sequence 12191	AL031295	Genbank	4376011
Sequence 12192	AL031297	Genbank	4902714
Sequence 12193	AL031427	Genbank	4835258
Sequence 12194	AL031432	Genbank	4375969
Sequence 12195	AL031665	Genbank	4826504
Sequence 12196	AL031681	Genbank	9408729
Sequence 12197	AL031718	Genbank	5805110
Sequence 12198	AL031774	Genbank	3881948
Sequence 12199	AL033377	Genbank	4826462
Sequence 12200	AL033397	Genbank	4902626
Sequence 12201	AL033517	Genbank	4140315
Sequence 12202	AL033522	Genbank	4191258
Sequence 12203	AL033531	Genbank	6807582
Sequence 12204	AL034548	Genbank	7263904
Sequence 12205	AL035209	Genbank	4160217
Sequence 12206	AL035249	Genbank	4688857
Sequence 12207	AL035398	Genbank	5262352
Sequence 12207	AL035411	Genbank	6136940
Sequence 12209	AL035413	Genbank	6010110
Sequence 12210	AL035454	Genbank	9368364
Sequence 12211	AL035458	Genbank	6624641
•	AL035461	Genbank	5123778
Sequence 12212		Genbank	9581764
Sequence 12213	AL035541	Genbank	
Sequence 12214	AL035610		6249454
Sequence 12215	AL035699	Genbank	4826515
Sequence 12216	AL049280	Genbank	4500037
Sequence 12217	AL049294	Genbank	4500055
Sequence 12218	AL049447	Genbank	4500230
Sequence 12219	AL049461	Genbank	4500250
Sequence 12220	AL049552	Genbank	6010194
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Sequence 12223	AL049694	Genbank	5650653
Sequence 12224	AL049748	Genbank	6572235
Sequence 12225	AL049766	Genbank	5763746
Sequence 12226	AL049779	Genbank	8176894
Sequence 12227	AL049829	Genbank	8217859
Sequence 12228	AL049844	Genbank	5924019
Sequence 12229	AL049849	Genbank	4826529
Sequence 12230	AL049869	Genbank	8217861
Sequence 12231	AL049870	Genbank	7248268
Sequence 12232	AL049911	Genbank	6114591
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Sequence 12234	AL049962	Genbank	4884212
Sequence 12235	AL049964	Genbank	4884214
Sequence 12236	AL049968	Genbank	4884217
Sequence 12237	AL050016	Genbank	4884262
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Sequence 12238	AL050089	Genbank	4884107
Sequence 12239	AL050275	Genbank	4886500
Sequence 12240	AL050277	Genbank	4886454
Sequence 12241	AL050343	Genbank	6137008
Sequence 12242	AL078463	Genbank	5777575
Sequence 12243	AL078621	Genbank	6013067
	AL079333	Genbank	5738434
Sequence 12244			
Sequence 12245	AL080080	Genbank	5262491
Sequence 12246	AL080102	Genbank	5262526
Sequence 12247	AL080193	Genbank	5262674
Sequence 12248	AL080276	Genbank	5763753
	AL109672	Genbank	5689836
Sequence 12249			
Sequence 12250	AL109759	Genbank	8176900
Sequence 12251	AL109798	Genbank	6453386
Sequence 12252	AL109865	Genbank	6911646
Sequence 12253	AL110183	Genbank	5817095
Sequence 12254	AL110185	Genbank	5817098
Sequence 12255	AL110193	Genbank	5817109
Sequence 12256	AL110194	Genbank	5817111
Sequence 12257	AL110202	Genbank	5817121
Sequence 12258	AL110271	Genbank	5817083
Sequence 12259	AL117350	Genbank	7159817
	AL117450	Genbank	5911900
Sequence 12260			
Sequence 12261	AL117461	Genbank	5911922
Sequence 12262	AL117482	Genbank	5911960
Sequence 12263	AL117648	Genbank	5912240
Sequence 12264	AL117662	Genbank	5912257
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Sequence 12266	AL121652	Genbank	7159615
	AL121716	Genbank	7159797
Sequence 12267			
Sequence 12268	AL121790	Genbank	8919824
Sequence 12269	AL121823	Genbank	6635879
Sequence 12270	AL121852	Genbank	8176919
Sequence 12271	AL121963	Genbank	7161783
Sequence 12272	AL122036	Genbank	6093222
Sequence 12273	AL122066	Genbank	6102860
	AL133010	Genbank	6453416
Sequence 12274			
Sequence 12275	AL133060	Genbank	6453483
Sequence 12276	AL133069	Genbank	6453507
Sequence 12277	AL133215	Genbank	7228177
Sequence 12278	AL133216	Genbank	6983473
Sequence 12279	AL133387	Genbank	7076389
Sequence 12280	AL133453	Genbank	7009594
		Genbank	6599181
Sequence 12281	AL133593		
Sequence 12282	AL133662	Genbank	6599256
Sequence 12283	AL135744	Genbank	6682291
Sequence 12284	AL135959	Genbank	6635874
Sequence 12285	AL136296	Genbank	8217913
Sequence 12286	AL136504	Genbank	7018294
Sequence 12287	AL136831	Genbank	6807666
		Genbank	6807703
Sequence 12288	AL137269		
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Sequence 12290	AL137454	Genbank	6808031
Sequence 12291	AL137481	Genbank	6808096
Sequence 12292	AL137542	Genbank	6808221
Sequence 12293	AL137673	Genbank	6807841
3545300 12200		#:#:::= #::	- · ·

Sequence 12294	AL137714	Genbank	6808128
Sequence 12295	AL137721	Genbank	6808159
Sequence 12296	AL137723	Genbank	6808203
Sequence 12297	AL137818	Genbank	7009598
Sequence 12298	AL157442	Genbank	7018558
Sequence 12299	AL157468	Genbank	7018484
Sequence 12300	AL157493	Genbank	7018539
Sequence 12301	AL158493	Genbank	7161641
Sequence 12302	AL158547	Genbank	7161742
Sequence 12303	AP000008	Genbank	3132318
Sequence 12304	AP000009	Genbank	4666255
Sequence 12305	AP000037	Genbank	3132347
Sequence 12306	AP000038	Genbank	3132348
Sequence 12307	AP000346	Genbank	5103009
Sequence 12308	AP000347	Genbank	5103010
Sequence 12309	AP000473	Genbank	6942329
Sequence 12310	AP000497	Genbank	5926684
Sequence 12311	AP000525	Genbank	5931503
Sequence 12312	AP000547	Genbank	5931525
Sequence 12313	AP000547 AP000692	Genbank	6693635
Sequence 12314	AP000962	Genbank	6942330
	AP000962 AP001063	Genbank	6693613
Sequence 12315	AP001003 AP001115	Genbank	6899854
Sequence 12316	AP001113	Genbank	
Sequence 12317	AP001136 AP001137	Genbank	6970360 6970361
Sequence 12318	AP001137 AP001138	Genbank	7262569
Sequence 12319			
Sequence 12320	AP001172	Genbank Genbank	6983884
Sequence 12321	AP001343		7209831
Sequence 12322	D00596	Genbank	220135
Sequence 12323	D13119	Genbank	285909
Sequence 12324	D13318	Genbank	286026
Sequence 12325	D13634	Genbank	285992
Sequence 12326	D14658	Genbank	285940
Sequence 12327	D14665 D14694	Genbank Genbank	6630617
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Sequence 133		7709	Genbank	3650435
Sequence 133			Genbank	3882148
Sequence 133			Genbank	3882154
Sequence 133			Genbank	3882168
Sequence 133			Genbank	3882272
Sequence 133	-		Genbank	3882294
Sequence 133			Genbank	3882314
Sequence 133			Genbank	3882318
Sequence 133			Genbank	3882322
Sequence 133			Genbank	3882340
Sequence 133			Genbank	6970469
Sequence 133			Genbank	4587126
Sequence 133			Genbank	4512311
Sequence 133	-	•	Genbank	4519939
Sequence 133		••	Genbank	3869126
Sequence 133			Genbank	3885367
Sequence 133			Genbank	6456471
Sequence 133			Genbank	4760601
Sequence 133			Genbank	6518494
Sequence 134	HOU ABUZ		•. • · · · · · · · · · · · · · · · · · ·	4240148
Sequence 134			Genbank	4240146
Sequence 134			Genbank	
Sequence 134	103 AB02		Genbank	4240194
Sequence 134	104 AB02		Genbank	4240198
Sequence 134	105 AB02		Genbank	4240226
Sequence 134			Genbank	4240238
Sequence 134	_		Genbank	4240262
Sequence 134			Genbank	4240284
Sequence 134			Genbank	4240296
Sequence 134			Genbank	4003379
Sequence 134	111 AB02		Genbank	4003388
Sequence 134			Genbank	4417209
Sequence 134	113 AB02	21470	Genbank	4586521

Sequence 13414	AB021663	Genbank	4996450
Sequence 13415		Genbank	4996607
Sequence 13416		Genbank	6942217
Sequence 13417		Genbank	4210446
Sequence 13418		Genbank	4589623
		Genbank	4589625
Sequence 13419			4589657
Sequence 13420		Genbank	
Sequence 13421		Genbank	6519222
Sequence 13422		Genbank	6016837
Sequence 13423	AB025355	Genbank	5103146
Sequence 13424		Genbank	6520072
Sequence 13425		Genbank	6172220
Sequence 13426		Genbank	5103045
Sequence 13427		Genbank	6552364
Sequence 13428		Genbank	5689530
Sequence 13429		Genbank	5295999
		Genbank	6730949
Sequence 13430		Genbank	5732186
Sequence 13431			
Sequence 13432		Genbank	6329748
Sequence 13433	3 AB032957	Genbank	6329818
Sequence 13434		Genbank	6382013
Sequence 13435	AB032969	Genbank	6329965
Sequence 13436		Genbank	6330242
Sequence 13437		Genbank	6330350
Sequence 13438		Genbank	6330364
Sequence 13439		Genbank	6330623
Sequence 13440		Genbank	6330797
Sequence 13441		Genbank	6330811
Sequence 13442		Genbank	6330846
		Genbank	6330860
Sequence 13443		Genbank	6382025
Sequence 13444		_	6331225
Sequence 13445		Genbank	
Sequence 13446		Genbank	6899845
Sequence 13447		Genbank	7243064
Sequence 13448	3 AB037767	Genbank	7243072
Sequence 13449	AB037807	Genbank	7243152
Sequence 13450	AB037813	Genbank	7243164
Sequence 1345	AC000015	Genbank	4263638
Sequence 13452		Genbank	4417318
Sequence 1345		Genbank	4731059
Sequence 13454		Genbank	4678240
Sequence 1345		Genbank	1809224
Sequence 1345		Genbank	6970735
		Genbank	2270906
Sequence 1345		Genbank	2133911
Sequence 1345	B AC000394		2133862
Sequence 1345	AC001226	Genbank	
Sequence 1346	O AC001234	Genbank	4028939
Sequence 1346		Genbank	2226439
Sequence 1346		Genbank	2342716
Sequence 1346		Genbank	2576343
Sequence 1346		Genbank	2076723
Sequence 1346	5 AC002081	Genbank	2078453
Sequence 1346		Genbank	2155224
Sequence 1346		Genbank	2599243
Sequence 1346		Genbank	2828783
Sequence 1346		Genbank	2961443
Sequence 1540	J 70002000	Johnann	2001-1-10

Sequence 13470	AC002365	Genbank	2358015
Sequence 13471	AC002366	Genbank	2739349
Sequence 13472	AC002394	Genbank	2815550
Sequence 13473	AC002395	Genbank	3540146
Sequence 13474	AC002398	Genbank	2529398
Sequence 13475	AC002401	Genbank	2599240
Sequence 13476	AC002487	Genbank	2341013
	AC002522	Genbank	7124014
Sequence 13477		Genbank	2393733
Sequence 13478	AC002542	Genbank	3337382
Sequence 13479	AC002544		2580474
Sequence 13480	AC002558	Genbank	
Sequence 13481	AC003004	Genbank	2522284
Sequence 13482	AC003007	Genbank	2911728
Sequence 13483	AC003026	Genbank	2815549
Sequence 13484	AC003029	Genbank	9965520
Sequence 13485	AC003037	Genbank	2920805
Sequence 13486	AC003041	Genbank	3264572
Sequence 13487	AC003046	Genbank	10043285
Sequence 13488	AC003070	Genbank	2865212
Sequence 13489	AC003103	Genbank	2842782
Sequence 13490	AC003104	Genbank	3253131
Sequence 13491	AC003663	Genbank	3097871
Sequence 13492	AC003682	Genbank	3264845
Sequence 13493	AC003688	Genbank	3789719
Sequence 13494	AC003963	Genbank	2995604
	AC003989	Genbank	2772538
Sequence 13495	AC003989 AC004006	Genbank	2781390
Sequence 13496		Genbank	2781380
Sequence 13497	AC004013	• • • • • • • • • • • • • • • • • • • •	2822136
Sequence 13498	AC004073	Genbank	
Sequence 13499	AC004084	Genbank	2822156
Sequence 13500	AC004106	Genbank	3150276
Sequence 13501	AC004126	Genbank	3242744
Sequence 13502	AC004142	Genbank	2880078
Sequence 13503	AC004148	Genbank	3482960
Sequence 13504	AC004152	Genbank	2894631
Sequence 13505	AC004156	Genbank	2896799
Sequence 13506	AC004158	Genbank	2982170
Sequence 13507	AC004221	Genbank	9665053
Sequence 13508	AC004222	Genbank	3108042
Sequence 13509	AC004223	Genbank	3253129
Sequence 13510	AC004227	Genbank	2911718
Sequence 13511	AC004228	Genbank	4263838
Sequence 13512	AC004233	Genbank	2914673
Sequence 13513	AC004236	Genbank	2914668
Sequence 13514	AC004258	Genbank	2924759
Sequence 13515	AC004263	Genbank	2935616
Sequence 13516	AC004381	Genbank	2982169
Sequence 13517	AC004382	Genbank	3252819
Sequence 13517	AC004388	Genbank	3046271
		Genbank	3873185
Sequence 13519	AC004408	Genbank	3617739
Sequence 13520	AC004466		
Sequence 13521	AC004477	Genbank	3688107
Sequence 13522	AC004494	Genbank	2996647
Sequence 13523	AC004507	Genbank	2996632
Sequence 13524	AC004520	Genbank	3004572
Sequence 13525	AC004522	Genbank	3006227

Sequence 13526	AC004527	Genbank	4760422
Sequence 13527	AC004531	Genbank	3337392
Sequence 13528	AC004539	Genbank	3041851
Sequence 13529	AC004582	Genbank	3337307
Sequence 13530	AC004584	Genbank	3417305
Sequence 13531	AC004585	Genbank	3212882
Sequence 13532	AC004587	Genbank	3150014
Sequence 13533	AC004600	Genbank	4388746
Sequence 13534	AC004601	Genbank	3395446
Sequence 13535	AC004611	Genbank	3080643
Sequence 13536	AC004617	Genbank	3738114
Sequence 13537	AC004622	Genbank	3090700
Sequence 13538	AC004634	Genbank	3094993
Sequence 13539	AC004677	Genbank	3126881
Sequence 13540	AC004686	Genbank	3688105
Sequence 13541	AC004760	Genbank	3168626
Sequence 13542	AC004765	Genbank	4731046
Sequence 13543	AC004770	Genbank	3212836
Sequence 13544	AC004798	Genbank	3184497
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Sequence 13546	AC004812	Genbank	3970970
Sequence 13547	AC004813	Genbank	4926912
Sequence 13548	AC004832	Genbank	6624129
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Sequence 13550	AC004842	Genbank	4753286
Sequence 13551	AC004846	Genbank	7243869
Sequence 13552	AC004854	Genbank	4827328
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Sequence 13554	AC004890	Genbank	4508146
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Sequence 13556	AC004908	Genbank	4156179
Sequence 13557	AC004938	Genbank	3213059
Sequence 13558	AC004941	Genbank	4454517
Sequence 13559	AC004950	Genbank	4753269
Sequence 13560	AC004955	Genbank	4753267
Sequence 13561	AC004969	Genbank	4156159
Sequence 13562	AC004982	Genbank	3419846
Sequence 13563	AC004983	Genbank	4309885
Sequence 13564	AC004985	Genbank	5708490
Sequence 13565	AC004987	Genbank	3213010
Sequence 13566	AC004990	Genbank	3924668
Sequence 13567	AC004999	Genbank	3970963
Sequence 13568	AC005005	Genbank	4156153
Sequence 13569	AC005007	Genbank	4156151
Sequence 13570	AC005021	Genbank	3694657
Sequence 13571	AC005031	Genbank	3688109
Sequence 13572	AC005037	Genbank	4827310
Sequence 13573	AC005041	Genbank	4508118
Sequence 13574	AC005042	Genbank	4156138
Sequence 13575	AC005046	Genbank	6094632
Sequence 13576	AC005076	Genbank	4508110
Sequence 13577	AC005082	Genbank	5836159
Sequence 13578	AC005084	Genbank	3659503
Sequence 13579	AC005088	Genbank	4753221
Sequence 13580	AC005095	Genbank	4753220
Sequence 13581	AC005104	Genbank	4218027

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Sequence 13582	AC005166	Genbank	3242769
Sequence 13583	AC005187	Genbank	3522927
Sequence 13584	AC005189	Genbank	3264580
Sequence 13585	AC005193	Genbank	3309101
			3273381
Sequence 13586	AC005203	Genbank	
Sequence 13587	AC005212	Genbank	3287444
Sequence 13588	AC005251	Genbank	3287718
Sequence 13589	AC005253	Genbank	3288886
Sequence 13590	AC005261	Genbank	3289984
Sequence 13591	AC005262	Genbank	3289981
	AC005300	Genbank	6492490
Sequence 13592			
Sequence 13593	AC005318	Genbank	3885345
Sequence 13594	AC005332	Genbank	3659494
Sequence 13595	AC005360	Genbank	3367514
Sequence 13596	AC005363	Genbank	3367511
Sequence 13597	AC005366	Genbank	3367508
Sequence 13598	AC005387	Genbank	3395417
Sequence 13599	AC005391	Genbank	3399673
Sequence 13600	AC005406	Genbank	4580403
Sequence 13601	AC005409	Genbank	4249432
Sequence 13602	AC005410	Genbank	3450753
Sequence 13603	AC005484	Genbank	5091654
Sequence 13604	AC005488	Genbank	5836194
Sequence 13605	AC005509	Genbank	4176355
Sequence 13606	AC005531	Genbank	4153875
		Genbank	3478668
Sequence 13607	AC005550		
Sequence 13608	AC005562	Genbank	4153858
Sequence 13609	AC005613	Genbank	3540152
Sequence 13610	AC005627	Genbank	7243867
Sequence 13611	AC005629	Genbank	7243877
Sequence 13612	AC005630	Genbank	4159882
Sequence 13613	AC005661	Genbank	3818348
Sequence 13614	AC005668	Genbank	3641771
Sequence 13615	AC005669	Genbank	3702356
		Genbank	3810672
Sequence 13616	AC005726		
Sequence 13617	AC005730	Genbank	3779042
Sequence 13618	AC005740	Genbank	3687210
Sequence 13619	AC005768	Genbank	6598827
Sequence 13620	AC005785	Genbank	3702290
Sequence 13621	AC005789	Genbank	3702281
Sequence 13622	AC005828	Genbank	3789713
Sequence 13623	AC005829	Genbank	3873300
		C. C	3849820
Sequence 13624	AC005837	Genbank	
Sequence 13625	AC005838	Genbank	3947427
Sequence 13626	AC005839	Genbank	4079626
Sequence 13627	AC005841	Genbank	4731044
Sequence 13628	AC005876	Genbank	6249672
Sequence 13629	AC005880	Genbank	6249667
Sequence 13630	AC005899	Genbank	3935221
Sequence 13631	AC005912	Genbank	4165005
		Genbank	4309927
Sequence 13632	AC005923		
Sequence 13633	AC005924	Genbank	4309926
Sequence 13634	AC005969	Genbank	4309946
Sequence 13635	AC005998	Genbank	5649387
Sequence 13636	AC006009	Genbank	4753278
Sequence 13637	AC006011	Genbank	4508143
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Sequence 13638	AC006017	Genbank	4508141
Sequence 13639	AC006023	Genbank	4753260
			6042101
Sequence 13640	AC006026	Genbank	
Sequence 13641	AC006027	Genbank	4156149
Sequence 13642	AC006050	Genbank	4079627
Sequence 13643	AC006057	Genbank	4731048
Sequence 13644	AC006064	Genbank	4572650
Sequence 13645	AC006084	Genbank	3953485
Sequence 13646	AC006088	Genbank	4204701
Sequence 13647	AC006129	Genbank	3970933
Sequence 13648	AC006137	Genbank	9887710
Sequence 13649	AC006141	Genbank	4580411
Sequence 13650	AC006153	Genbank	4309917
Sequence 13651	AC006160	Genbank	5701616
Sequence 13652	AC006205	Genbank	5815508
Sequence 13653	AC006206	Genbank	4309925
Sequence 13654	AC006207	Genbank	4309924
Sequence 13655	AC006238	Genbank	4204704
Sequence 13656	AC006251	Genbank	4309945
		Genbank	4726099
Sequence 13657	AC006254		
Sequence 13658	AC006255	Genbank	5306223
Sequence 13659	AC006257	Genbank	4092478
Sequence 13660	AC006273	Genbank	4096060
Sequence 13661	AC006347	Genbank	4309919
Sequence 13662	AC006349	Genbank	7243873
	AC006409	Genbank	4454486
Sequence 13663			
Sequence 13664	AC006430	Genbank	6453929
Sequence 13665	AC006449	Genbank	6102666
Sequence 13666	AC006450	Genbank	6997336
Sequence 13667	AC006475	Genbank	4753283
Sequence 13668	AC006480	Genbank	6289252
Sequence 13669	AC006487	Genbank	4314418
	AC006504	Genbank	4220442
Sequence 13670			
Sequence 13671	AC006509	Genbank	5815507
Sequence 13672	AC006512	Genbank	4926863
Sequence 13673	AC006518	Genbank	4713939
Sequence 13674	AC006536	Genbank	4235135
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Sequence 13676	AC006561	Genbank	4558534
	AC006581	Genbank	4914350
Sequence 13677	_		
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Sequence 13679	AC006930	Genbank	4309680
Sequence 13680	AC006973	Genbank	4699968
Sequence 13681	AC006987	Genbank	5123989
Sequence 13682	AC007014	Genbank	4371263
Sequence 13683	AC007029	Genbank	7243872
		Genbank	5708475
Sequence 13684	AC007038		
Sequence 13685	AC007041	Genbank	5708471
Sequence 13686	AC007050	Genbank	6456174
Sequence 13687	AC007055	Genbank	4885691
Sequence 13688	AC007057	Genbank	4680440
Sequence 13689	AC007114	Genbank	4581193
Sequence 13690	AC007130	Genbank	5001538
Sequence 13691	AC007172	Genbank	4731066
Sequence 13692	AC007204	Genbank	4559317
Sequence 13693	AC007207	Genbank	6466489

Sequence 13694	AC007216	Genbank	6806842
Sequence 13695	AC007225	Genbank	6715703
Sequence 13696	AC007229	Genbank	4567173
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Sequence 13697	AC007239	Genbank	5523828
Sequence 13698	AC007252	Genbank	5091646
Sequence 13699	AC007314	Genbank	4662678
Sequence 13700	AC007324	Genbank	7923342
			4587630
Sequence 13701	AC007327	Genbank	
Sequence 13702	AC007390	Genbank	6358845
Sequence 13703	AC007393	Genbank	5306300
Sequence 13704	AC007446	Genbank	5069495
Sequence 13705	AC007528	Genbank	5263309
Sequence 13706	AC007536	Genbank	5917929
			4982536
Sequence 13707	AC007541	Genbank	
Sequence 13708	AC007546	Genbank	5668756
Sequence 13709	AC007551	Genbank	4809347
Sequence 13710	AC007566	Genbank	4835815
Sequence 13711	AC007766	Genbank	5030437
Sequence 13712	AC007848	Genbank	5923645
			5649179
Sequence 13713	AC007860	Genbank	
Sequence 13714	AC007868	Genbank	6041758
Sequence 13715	AC007934	Genbank	6721208
Sequence 13716	AC008009	Genbank	5668757
Sequence 13717	AC008018	Genbank	7958973
Sequence 13718	AC008102	Genbank	9797815
Sequence 13719	AC008115	Genbank	5801654
			6006046
Sequence 13720	AC008123	Genbank	
Sequence 13721	AC008151	Genbank	5629925
Sequence 13722	AC008498	Genbank	6850299
Sequence 13723	AC008545	Genbank	6721145
Sequence 13724	AC008583	Genbank	7158894
Sequence 13725	AC008893	Genbank	7259691
	AC008981	Genbank	6721142
Sequence 13726			7243882
Sequence 13727	AC009298	Genbank	
Sequence 13728	AC009363	Genbank	6648145
Sequence 13729	AC009509	Genbank	6492473
Sequence 13730	AC009946	Genbank	6604542
Sequence 13731	AC010077	Genbank	5870275
Sequence 13732	AC010168	Genbank	6855156
	AC010197	Genbank	6143841
Sequence 13733		Genbank	
Sequence 13734	AC010436	-	6938861
Sequence 13735	AC010525	Genbank	7248932
Sequence 13736	AC010582	Genbank	6721135
Sequence 13737	AC010722	Genbank	6587939
Sequence 13738	AC011331	Genbank	9929686
Sequence 13739	AC011599	Genbank	7113871
	AC012085	Genbank	6634755
Sequence 13740			
Sequence 13741	AC012087	Genbank	7114465
Sequence 13742	AC016831	Genbank	6539289
Sequence 13743	AC016941	Genbank	8567654
Sequence 13744	AC020663	Genbank	6682593
Sequence 13745	AC021049	Genbank	7248920
Sequence 13746	AF000982	Genbank	2580549
Sequence 13747	AF001549	Genbank	3355302
		Genbank	2339976
Sequence 13748	AF001552		
Sequence 13749	AF002668	Genbank	2232173

Sequence 13750	AF004162	Genbank	3046385
Sequence 13751	AF004337	Genbank	2245557
Sequence 13752	AF004338	Genbank	2245558
Sequence 13753	AF004339	Genbank	2245559
	AF004342	Genbank	2245565
Sequence 13754		Genbank	2645428
Sequence 13755	AF006513		7629276
Sequence 13756	AF006621	Genbank	
Sequence 13757	AF007134	Genbank	2852608
Sequence 13758		Genbank	2852624
Sequence 13759	AF007189	Genbank	2459927
Sequence 13760	AF007544	Genbank	2970122
Sequence 13761	AF007872	Genbank	2358280
Sequence 13762	AF007875	Genbank	2258417
Sequence 13763		Genbank	2580619
Sequence 13764		Genbank	2338289
Sequence 13765		Genbank	6468761
		Genbank	2335186
Sequence 13766		Genbank	2343184
Sequence 13767	· ·	Genbank	3046899
Sequence 13768			2801421
Sequence 13769		Genbank	
Sequence 13770		Genbank	8148061
Sequence 13771	AF015812	Genbank	2599359
Sequence 13772	AF017790	Genbank	2501872
Sequence 13773	AF020352	Genbank	2655054
Sequence 13774	AF020782	Genbank	2522480
Sequence 13775		Genbank	2460317
Sequence 13776		Genbank	2465723
Sequence 13777		Genbank	2935182
Sequence 13778		Genbank	5712085
Sequence 13779		Genbank	3599965
Sequence 13780		Genbank	2598967
Sequence 13781		Genbank	3287488
		Genbank	2587057
Sequence 13782		Genbank	3403166
Sequence 13783	AF029786		4894208
Sequence 13784	AF031379	Genbank	2661038
Sequence 13785		Genbank	
Sequence 13786		Genbank	2661052
Sequence 13787		Genbank	2661064
Sequence 13788	3 AF035840	Genbank	3800739
Sequence 13789	AF036715	Genbank	4007385
Sequence 13790		Genbank	2708629
Sequence 13791		Genbank	6466790
Sequence 13792		Genbank	2795921
Sequence 13793		Genbank	3523122
Sequence 13794		Genbank	2828109
Sequence 13795		Genbank	2773157
Sequence 13796		Genbank	3366667
		Genbank	3869269
Sequence 13797		Genbank	3776241
Sequence 13798		Genbank	3169541
Sequence 13799			
Sequence 13800		Genbank	5326827
Sequence 1380		Genbank	4164445
Sequence 13802		Genbank	3348136
Sequence 13803		Genbank	2865655
Sequence 13804	4 AF047184	Genbank	2909859
Sequence 1380		Genbank	2909861
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<u>Table 3-5</u>

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Sequence 13806	AF047438	Genbank	3335131
Sequence 13807	AF047440	Genbank	3335135
Sequence 13808	AF047472	Genbank	2921872
Sequence 13809	AF050638	Genbank	5326819
Sequence 13810	AF050639	Genbank	4164453
Sequence 13811	AF052092	Genbank	3360398
Sequence 13812	AF052129	Genbank	3360438
	AF052164	Genbank	3360475
Sequence 13813	AF052182	Genbank	3360494
Sequence 13814		Genbank	3510694
Sequence 13815	AF052573		2967847
Sequence 13816	AF052578	Genbank	3135305
Sequence 13817	AF053356	Genbank	
Sequence 13818	AF054183	Genbank	4092053
Sequence 13819	AF054990	Genbank	3005703
Sequence 13820	AF055066	Genbank	3273727
Sequence 13821	AF055474	Genbank	nil
Sequence 13822	AF055584	Genbank	3649607
Sequence 13823	AF057356	Genbank	3063652
Sequence 13824	AF057735	Genbank	4165042
Sequence 13825	AF058293	Genbank	3047377
Sequence 13826	AF058988	Genbank	3300089
Sequence 13827	AF059527	Genbank	4454661
Sequence 13828	AF060798	Genbank	3372665
Sequence 13829	AF061737	Genbank	4335938
Sequence 13830	AF062077	Genbank	3641524
Sequence 13831	AF064019	Genbank	3347856
Sequence 13832	AF064861	Genbank	3171154
Sequence 13833	AF067139	Genbank	3337440
Sequence 13834	AF068302	Genbank	4584876
	AF068754	Genbank	3283408
Sequence 13835	AF068755	Genbank	4321979
Sequence 13836	AF069378	Genbank	3982733
Sequence 13837	AF069532	Genbank	3892190
Sequence 13838	AF070523	Genbank	3764088
Sequence 13839	AF070523 AF070539	Genbank	3387896
Sequence 13840		Genbank	3387921
Sequence 13841	AF070556	Genbank	3283898
Sequence 13842	AF070630		3283905
Sequence 13843	AF070635	Genbank	3283909
Sequence 13844	AF070638	Genbank	3283920
Sequence 13845	AF070646	Genbank	
Sequence 13846	AF070649	Genbank	3283923 4454685
Sequence 13847	AF070655	Genbank	
Sequence 13848	AF070659	Genbank	4454693
Sequence 13849	AF070669	Genbank	4454713
Sequence 13850	AF070673	Genbank	3978241
Sequence 13851	AF072097	Genbank	5725511
Sequence 13852	AF073298	Genbank	3641537
Sequence 13853	AF075061	Genbank	3377602
Sequence 13854	AF077030	Genbank	4689107
Sequence 13855	AF077036	Genbank	4689119
Sequence 13856	AF077043	Genbank	4689133
Sequence 13857		Genbank	4689135
Sequence 13858		Genbank	4689151
Sequence 13859		Genbank	4679013
Sequence 13860		Genbank	4679017
Sequence 13861	AF078847	Genbank	5531808

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Sequence 13862	AF078850	Genbank	5531814
Sequence 13863	AF082283	Genbank	4092066
Sequence 13864	AF082513	Genbank	3649656
Sequence 13865	AF083441	Genbank	5813822
Sequence 13866	AF085844	Genbank	3483158
	AF086210	Genbank	3483555
Sequence 13867			3483674
Sequence 13868	AF086329	Genbank	
Sequence 13869	AF086337	Genbank	3483682
Sequence 13870	AF086378	Genbank	3483723
Sequence 13871	AF086490	Genbank	3483835
Sequence 13872	AF086557	Genbank	3483902
Sequence 13873	AF086564	Genbank	3483909
Sequence 13874	AF086628	Genbank	4240457
Sequence 13875	AF087020	Genbank	3851144
Sequence 13876	AF087160	Genbank	3869380
Sequence 13877	AF087959	Genbank	3523165
	AF088060	Genbank	3523266
Sequence 13878	AF088071	Genbank	3523277
Sequence 13879		Genbank	4063630
Sequence 13880	AF090095		6690255
Sequence 13881	AF090947	Genbank	
Sequence 13882	AF091075	Genbank	3859987
Sequence 13883	AF092094	Genbank	4426606
Sequence 13884	AF092124	Genbank	3659900
Sequence 13885	AF092128	Genbank	5138905
Sequence 13886	AF092130	Genbank	5138909
Sequence 13887	AF092135	Genbank	5138919
Sequence 13888	AF097026	Genbank	6409120
Sequence 13889	AF097514	Genbank	4808600
Sequence 13890	AF098162	Genbank	3929582
Sequence 13891	AF100615	Genbank	4808624
Sequence 13892	AF100616	Genbank	4808626
Coguence 13032	AF100748	Genbank	5410281
Sequence 13893	AF100757	Genbank	5410299
Sequence 13894	AF100757	Genbank	5410305
Sequence 13895		Genbank	4092759
Sequence 13896	AF102803		5880481
Sequence 13897		Genbank	
Sequence 13898	AF104913	Genbank	3941723
Sequence 13899	AF106622	Genbank	4378528
Sequence 13900	AF106681	Genbank	5410327
Sequence 13901	AF109134	Genbank	4139227
Sequence 13902	AF109718	Genbank	3978547
Sequence 13903	AF110460	Genbank	4324698
Sequence 13904	AF111847	Genbank	7211441
Sequence 13905	AF112213	Genbank	6563213
Sequence 13906	AF112222	Genbank	6563229
Sequence 13907	AF112227	Genbank	4545218
Sequence 13908	AF113011	Genbank	6642745
Sequence 13909	AF113016	Genbank	6642755
Sequence 13910	AF113141	Genbank	4512031
Sequence 13911	AF113535	Genbank	6523824
	AF113535 AF113540	Genbank	6523834
Sequence 13912		Genbank	5106955
Sequence 13913	AF113615		6855605
Sequence 13914	AF113680	Genbank	
Sequence 13915	AF113690	Genbank	6855619
Sequence 13916	AF113700	Genbank	6855634
Sequence 13917	AF113701	Genbank	6855635

	4545400	Cambank	4550272
Sequence 13918	AF115402	Genbank	4559272
Sequence 13919	AF116910	Genbank	4768837
Sequence 13920	AF117235	Genbank	6563243
Sequence 13921	AF117888	Genbank	5732617
Sequence 13922	AF118084	Genbank	6650813
Sequence 13923	AF118124	Genbank	4235636
Sequence 13924	AF118649	Genbank	6840946
Sequence 13925	AF119297	Genbank	4633508
	AF123052	Genbank	5106556
Sequence 13926	AF125530	Genbank	6563259
Sequence 13927		Genbank	6563275
Sequence 13928			6318547
Sequence 13929	AF126782	Genbank	7019596
Sequence 13930	AF127577	Genbank	
Sequence 13931	AF128527	Genbank	4928043
Sequence 13932	AF128536	Genbank	5305705
Sequence 13933	AF129756	Genbank	4337095
Sequence 13934	AF131738	Genbank	4406548
Sequence 13935	AF131748	Genbank	4406563
Sequence 13936		Genbank	4406576
Sequence 13937		Genbank	4406595
Sequence 13938		Genbank	4406602
Sequence 13939		Genbank	4406608
		Genbank	4406648
Sequence 13940		Genbank	4406677
Sequence 13941		- ·	4406702
Sequence 13942		Genbank	
Sequence 13943		Genbank	4680648
Sequence 13944		Genbank	4680654
Sequence 13945	AF132973	Genbank	4680716
Sequence 13946	AF132984	Genbank	4959567
Sequence 13947	AF134726	Genbank	4529886
Sequence 13948		Genbank	7259481
Sequence 13949	AF135488	Genbank	5668619
Sequence 13950		Genbank	5020073
Sequence 13951		Genbank	4761681
Sequence 13952	AF147378	Genbank	4761729
Sequence 13953		Genbank	4761763
Sequence 13954		Genbank	5107165
Sequence 13955		Genbank	7106825
		Genbank	5758136
Sequence 13956		Genbank	6166337
Sequence 13957		Genbank	4929554
Sequence 13958			4929566
Sequence 13959		Genbank	
Sequence 13960) AF151832	Genbank	4929616
Sequence 13961		Genbank	4929622
Sequence 13962		Genbank	4929708
Sequence 13963		Genbank	4929758
Sequence 13964	4 AF151905	Genbank	4929762
Sequence 13965	5 AF151906	Genbank	4929764
Sequence 13966		Genbank	5669134
Sequence 13967		Genbank	5932030
Sequence 13968		Genbank	5932031
Sequence 13969		Genbank	5231140
Sequence 13970		Genbank	5731112
Sequence 1397		Genbank	6959307
		Genbank	5714635
Sequence 13972		Genbank	6841181
Sequence 1397) ME101304	Germank	0041101

<u>Table 3-5</u>

Sequence 13974	AF161389	Genbank	6841191
Sequence 13975	AF161428	Genbank	6841269
Sequence 13976	AF161448	Genbank	6841309
	AF161464	Genbank	6841451
Sequence 13977			
Sequence 13978	AF161479	Genbank	6841481
Sequence 13979	AF161481	Genbank	6841485
Sequence 13980	AF161485	Genbank	6841493
Sequence 13981	AF161500	Genbank	6841523
Sequence 13982	AF161503	Genbank	6841529
Sequence 13983	AF161509	Genbank	6841541
Sequence 13984	AF161512	Genbank	6841547
	AF161541	Genbank	6841349
Sequence 13985			
Sequence 13986	AF161556	Genbank	6841379
Sequence 13987	AF165926	Genbank	5817858
Sequence 13988	AF168956	Genbank	5702387
Sequence 13989	AF172066	Genbank	5733811
Sequence 13990	AF176315	Genbank	5823349
Sequence 13991	AF181896	Genbank	6272683
Sequence 13992	AF187320	Genbank	6164847
	AF187554	Genbank	6653225
Sequence 13993			6425043
Sequence 13994	AF188745	Genbank	
Sequence 13995	AF189712	Genbank	6760474
Sequence 13996	AF190465	Genbank	6120105
Sequence 13997	AF191018	Genbank	6457339
Sequence 13998	AF191298	Genbank	7656642
Sequence 13999	AF201077	Genbank	6456748
Sequence 14000	AF203815	Genbank	6979641
Sequence 14001	AF205588	Genbank	6531675
	AF207550	Genbank	6470333
Sequence 14002		Genbank	6942010
Sequence 14003	AF216305		7158868
Sequence 14004	AF228703	Genbank	
Sequence 14005	AF228704	Genbank	7158871
Sequence 14006	AJ000519	Genbank	2739214
Sequence 14007	AJ001014	Genbank	3171909
Sequence 14008	AJ001866	Genbank	2969902
Sequence 14009	AJ001981	Genbank	2546978
Sequence 14010	AJ002030	Genbank	2570006
Sequence 14011	AJ002308	Genbank	2959871
	AJ002385	Genbank	2597930
Sequence 14012			3687321
Sequence 14013	AJ006470	Genbank	
Sequence 14014	AJ010069	Genbank	3483012
Sequence 14015	AJ010597	Genbank	3559873
Sequence 14016	AJ012409	Genbank	3881975
Sequence 14017	AJ012499	Genbank	5441359
Sequence 14018	AJ223812	Genbank	2894518
Sequence 14019	AJ224442	Genbank	2911586
Sequence 14020	AJ230595	Genbank	4583774
Sequence 14021	AJ230596	Genbank	4583775
	AJ230605	Genbank	4583784
Sequence 14022			4826530
Sequence 14023	AJ238243	Genbank	
Sequence 14024	AJ238283	Genbank	4704219
Sequence 14025	AJ238374	Genbank	6634418
Sequence 14026	AJ245719	Genbank	6688168
Sequence 14027	AJ249731	Genbank	5921469
Sequence 14028		Genbank	6013005
Sequence 14029		Genbank	6996445
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<u>Table 3-5</u>

Sequence 14030	AJ251961	Genbank	6634453
Sequence 14031	AJ271216	Genbank	6966966
Sequence 14032	AK000018	Genbank	7019828
Sequence 14033	AK000023	Genbank	7019837
Sequence 14034	AK000035	Genbank	7019855
	AK000208	Genbank	7020141
Sequence 14035		Genbank	7020274
Sequence 14036	AK000291	Genbank	7020274
Sequence 14037	AK000348	· · · · · · · · · · · · · · · · · · ·	
Sequence 14038	AK000365	Genbank	7020402
Sequence 14039	AK000395	Genbank	7020455
Sequence 14040	AK000426	Genbank	7020505
Sequence 14041	AK000434	Genbank	7020520
Sequence 14042	AK000453	Genbank	7020550
Sequence 14043	AK000462	Genbank	7020566
Sequence 14044	AK000470	Genbank	7020580
Sequence 14045	AK000507	Genbank	7020644
Sequence 14046	AK000548	Genbank	7020717
Sequence 14047	AK000560	Genbank	7020738
Sequence 14048	AK000571	Genbank	7020756
Sequence 14049	AK000585	Genbank	7020779
Sequence 14050	AK000615	Genbank	7020827
Sequence 14051	AK000620	Genbank	7020834
Sequence 14052	AK000624	Genbank	7020839
	AK000655	Genbank	7020888
Sequence 14053 Sequence 14054	AK000655 AK000667	Genbank	7020906
	AK000669	Genbank	7020909
Sequence 14055	AK000009 AK000721	Genbank	7020985
Sequence 14056		Genbank	7020990
Sequence 14057	AK000725	Genbank	7020990
Sequence 14058	AK000739		7021014
Sequence 14059	AK000745	Genbank	
Sequence 14060	AK000752	Genbank	7021035
Sequence 14061	AK000765	Genbank	7021058
Sequence 14062	AK000800	Genbank	7021104
Sequence 14063	AK000818	Genbank	7021128
Sequence 14064	AK000826	Genbank	7021139
Sequence 14065	AK000866	Genbank	7021190
Sequence 14066	AK000891	Genbank	7021840
Sequence 14067	AK000933	Genbank	7021908
Sequence 14068	AK000934	Genbank	7021909
Sequence 14069	AK000959	Genbank	7021945
Sequence 14070	AK000991	Genbank	7021994
Sequence 14071	AK001006	Genbank	7022017
Sequence 14072	AK001093	Genbank	7022142
Sequence 14073	AK001105	Genbank	7022160
Sequence 14074	AK001154	Genbank	7022228
Sequence 14075	AK001237	Genbank	7022366
Sequence 14076	AK001285	Genbank	7022444
Sequence 14077	AK001330	Genbank	7022520
Sequence 14078	AK001387	Genbank	7022615
Sequence 14079	AK001413	Genbank	7022655
Sequence 14080	AK001441	Genbank	7022699
Sequence 14081	AK001451	Genbank	7022717
Sequence 14082	AK001469	Genbank	7022747
Sequence 14083	AK001493	Genbank	7022783
Sequence 14084	AK001536	Genbank	7022851
Sequence 14085	AK001571	Genbank	7022906
2042000000			

Sequence 14086	AK001656	Genbank	7023047
Sequence 14087	AK001690	Genbank	7023105
Sequence 14088	AK001691	Genbank	7023106
Sequence 14089	AK001717	Genbank	7023151
Sequence 14090	AK001718	Genbank	7023153
Sequence 14091	AK001747	Genbank	7023203
Sequence 14092	AK001762	Genbank	7023231
Sequence 14093	AK001855	Genbank	7023382
Sequence 14094	AK001833	Genbank	7023416
Sequence 14094 Sequence 14095	AK001877 AK001878	Genbank	7023418
	AK001978 AK001961	Genbank	7023549
Sequence 14096	AK001961 AK002002	Genbank	7023620
Sequence 14097		Genbank	7023689
Sequence 14098	AK002046	Genbank	7023781
Sequence 14099	AK002102	Genbank	7023781
Sequence 14100	AK002169		7023004
Sequence 14101	AK002190	Genbank	
Sequence 14102	AL008582	Genbank	5360981
Sequence 14103	AL008639	Genbank	6599068
Sequence 14104	AL008721	Genbank	3171883
Sequence 14105	AL008725	Genbank	2791551
Sequence 14106	AL009051	Genbank	2995195
Sequence 14107	AL009181	Genbank	2853179
Sequence 14108	AL020993	Genbank	3980107
Sequence 14109	AL020995	Genbank	5881341
Sequence 14110	AL021069	Genbank	2853183
Sequence 14111	AL021368	Genbank	3080468
Sequence 14112	AL021546	Genbank	2826890
Sequence 14113	AL021920	Genbank	3264535
Sequence 14114	AL022165	Genbank	3281985
Sequence 14115	AL022238	Genbank	4176442
Sequence 14116	AL022240	Genbank	4826471
Sequence 14117	AL022302	Genbank	5262844
Sequence 14118	AL022312	Genbank	4914501
Sequence 14119	AL022313	Genbank	4200326
Sequence 14120	AL022316	Genbank	4691242
Sequence 14121	AL022320	Genbank	4914513
Sequence 14122	AL022328	Genbank	5263010
Sequence 14123	AL022394	Genbank	9581758
Sequence 14124	AL022395	Genbank	4468287
Sequence 14125	AL022725	Genbank	5679748
Sequence 14126	AL023284	Genbank	3355875
Sequence 14127	AL023494	Genbank	5360992
Sequence 14128	AL023798	Genbank	3550031
Sequence 14129	AL024493	Genbank	3288439
Sequence 14130	AL030996	Genbank	3688349
Sequence 14131	AL031003	Genbank	4007185
Sequence 14132	AL031259	Genbank	3790132
Sequence 14133	AL031274	Genbank	3421083
	AL031287	Genbank	4914533
Sequence 14134 Sequence 14135	AL031297 AL031295	Genbank	4376011
	AL031293 AL031368	Genbank	4455593
Sequence 14136		Genbank	5002609
Sequence 14137	AL031390	Genbank	5931719
Sequence 14138	AL031428		4938290
Sequence 14139	AL031431	Genbank	4375969
Sequence 14140	AL031432	Genbank	4826442
Sequence 14141	AL031433	Genbank	4020442

Sequence 14142	AL031447	Genbank	4826431
Sequence 14143	AL031597	Genbank	4775640
Sequence 14144	AL031662	Genbank	9716901
Sequence 14145	AL031667	Genbank	9408727
	AL031681	Genbank	9408729
Sequence 14146		Genbank	9798435
Sequence 14147	AL031710		
Sequence 14148	AL031716	Genbank	4826474
Sequence 14149	AL031732	Genbank	4581343
Sequence 14150	AL031781	Genbank	4038570
Sequence 14151	AL032821	Genbank	4572584
Sequence 14152	AL033377	Genbank	4826462
Sequence 14153	AL033397	Genbank	4902626
Sequence 14154	AL033522	Genbank	4191258
Sequence 14155	AL033531	Genbank	6807582
Sequence 14156	AL033543	Genbank	4581346
Sequence 14157	AL034417	Genbank	5102616
•	AL034548	Genbank	7263904
Sequence 14158		Genbank	4160217
Sequence 14159	AL035209		6010110
Sequence 14160	AL035413	Genbank	
Sequence 14161	AL035420	Genbank	5918161
Sequence 14162	AL035425	Genbank	6465823
Sequence 14163	AL035458	Genbank	6624641
Sequence 14164	AL035541	Genbank	9581764
Sequence 14165	AL035652	Genbank	9650702
Sequence 14166	AL035681	Genbank	4902689
Sequence 14167	AL035699	Genbank	4826515
Sequence 14168	AL049249	Genbank	4499987
Sequence 14169	AL049282	Genbank	4500041
Sequence 14170	AL049404	Genbank	4500192
Sequence 14171	AL049557	Genbank	5596765
Sequence 14172	AL049563	Genbank	4902753
Sequence 14172	AL049564	Genbank	4902757
Sequence 14174	AL049589	Genbank	5679567
	AL049696	Genbank	5931893
Sequence 14175	AL049090 AL049742	Genbank	5419783
Sequence 14176		Genbank	5738437
Sequence 14177	AL049761		5763746
Sequence 14178	AL049766	Genbank	
Sequence 14179	AL049776	Genbank	6681704
Sequence 14180	AL049779	Genbank	8176894
Sequence 14181	AL049829	Genbank	8217859
Sequence 14182	AL049844	Genbank	5924019
Sequence 14183	AL049869	Genbank	8217861
Sequence 14184	AL049929	Genbank	4884174
Sequence 14185	AL049968	Genbank	4884217
Sequence 14186	AL049987	Genbank	4884238
Sequence 14187	AL050021	Genbank	4884264
Sequence 14188	AL050027	Genbank	4884266
Sequence 14189	AL050035	Genbank	4884276
Sequence 14190	AL050083	Genbank	4884317
Sequence 14191	AL050161	Genbank	4884375
	AL050101	Genbank	4884165
Sequence 14192			4884454
Sequence 14193	AL050215	Genbank	
Sequence 14194	AL050273	Genbank	4886450
Sequence 14195	AL050275	Genbank	4886500
Sequence 14196	AL050290	Genbank	4886512
Sequence 14197	AL050343	Genbank	6137008

Sequence 14198	AL050363	Genbank	4914597
Sequence 14199	AL050383	Genbank	4914585
Sequence 14200	AL050392	Genbank	4914613
Sequence 14201	AL078591	Genbank	6562085
	AL078621	Genbank	6013067
Sequence 14202			
Sequence 14203	AL079295	Genbank	5102607
Sequence 14204	AL079304	Genbank	8176897
Sequence 14205	AL079342	Genbank	6018784
Sequence 14206	AL080088	Genbank	5262504
Sequence 14207	AL080089	Genbank	5262506
Sequence 14208	AL080097	Genbank	5262519
Sequence 14209	AL080109	Genbank	5262535
Sequence 14210	AL080135	Genbank	5262576
Sequence 14211	AL080144	Genbank	5262592
Sequence 14212	AL080186	Genbank	5262664
Sequence 14213	AL080241	Genbank	5730204
Sequence 14214	AL080250	Genbank	5804917
		Genbank	5763753
Sequence 14215	AL080276		5918011
Sequence 14216	AL096773	Genbank	
Sequence 14217	AL096803	Genbank	6010162
Sequence 14218	AL109657	Genbank	6136991
Sequence 14219	AL109684	Genbank	5689805
Sequence 14220	AL109925	Genbank	7105935
Sequence 14221	AL110183	Genbank	5817095
Sequence 14222	AL110185	Genbank	5817098
Sequence 14223	AL110193	Genbank	5817109
Sequence 14224	AL110194	Genbank	5817111
Sequence 14225	AL110202	Genbank	5817121
Sequence 14226	AL110208	Genbank	5817127
Sequence 14227	AL110262	Genbank	5817224
Sequence 14228	AL117334	Genbank	7248316
Sequence 14229	AL117424	Genbank	5911856
Sequence 14230	AL117430	Genbank	5911865
Sequence 14231	AL117442	Genbank	5911885
	AL117442 AL117558	Genbank	5912097
Sequence 14232	AL117599	Genbank	5912167
Sequence 14233			
Sequence 14234	AL117621	Genbank	5912202
Sequence 14235	AL117648	Genbank	5912240
Sequence 14236	AL117662	Genbank	5912257
Sequence 14237	AL117666	Genbank	5912264
Sequence 14238	AL121576	Genbank	8176910
Sequence 14239	AL121580	Genbank	6456854
Sequence 14240	AL121603	Genbank	6434634
Sequence 14241	AL121656	Genbank	7159617
Sequence 14242	AL121782	Genbank	6456831
Sequence 14243	AL121790	Genbank	8919824
Sequence 14244	AL121913	Genbank	7161781
Sequence 14245	AL121963	Genbank	7161783
Sequence 14246	AL122003	Genbank	6580480
Sequence 14247	AL122013	Genbank	7242500
Sequence 14247	AL122010	Genbank	6093244
Sequence 14249	AL122066	Genbank	6102860
		Genbank	6102899
Sequence 14250	AL122089		
Sequence 14251	AL133051	Genbank	6453468
Sequence 14252	AL133060	Genbank	6453483
Sequence 14253	AL133094	Genbank	6453548

Sequence 14254	AL133238	Genbank	6624577
Sequence 14255	AL133241	Genbank	7630038
Sequence 14256	AL133243	Genbank	6491713
Sequence 14257	AL133246	Genbank	7594587
Sequence 14258	AL133370	Genbank	8248722
Sequence 14259	AL133453	Genbank	7009594
Sequence 14260	AL133455	Genbank	7019734
Sequence 14261	AL133555	Genbank	6599122
Sequence 14262	AL135744	Genbank	6682291
Sequence 14263	AL135858	Genbank	7159622
Sequence 14264	AL136296	Genbank	8217913
Sequence 14265	AL136504	Genbank	7018294
Sequence 14266	AL137440	Genbank	6808001
Sequence 14267	AL137450	Genbank	6808024
Sequence 14268	AL137489	Genbank	6808110
Sequence 14269	AL137496	Genbank	6808137
Sequence 14270	AL137553	Genbank	6808238
Sequence 14271	AL137567	Genbank	6808276
Sequence 14272	AL137659	Genbank	6807778
Sequence 14273	AL137673	Genbank	6807841
Sequence 14274	AL137681	Genbank	6807931
Sequence 14275	AL137714	Genbank	6808128
Sequence 14276	AL137750	Genbank	6808386
Sequence 14277	AL157428	Genbank	7018458
Sequence 14278	AL157464	Genbank	7018479
Sequence 14279	AP000008	Genbank	3132318
Sequence 14280	AP000031	Genbank	3132341
Sequence 14281	AP000038	Genbank	3132348
Sequence 14282	AP000351	Genbank	6069325
Sequence 14283	AP000354	Genbank	5103017
Sequence 14284	AP000473	Genbank	6942329
Sequence 14285	AP000525	Genbank	5931503
Sequence 14286	AP000526	Genbank	5931504
Sequence 14287	AP000962	Genbank	6942330
Sequence 14288	AP001055	Genbank	6693605
Sequence 14289	AP001136	Genbank	6970360
Sequence 14290	AP001137	Genbank	6970361
Sequence 14291	AP001172	Genbank	6983884
Sequence 14292	AP001343	Genbank	7209831
Sequence 14293	AP001346	Genbank	7209834
Sequence 14294	D10493	Genbank	219932
Sequence 14295	D12485	Genbank	219943
Sequence 14296	D14082	Genbank	468933
Sequence 14297	D14533	Genbank	286028
Sequence 14298	D14540	Genbank	467511
Sequence 14299	D14658	Genbank	285940
Sequence 14300	D14665	Genbank	6630617
Sequence 14301	D14696	Genbank	285962
Sequence 14302	D14710	Genbank	559324
Sequence 14303	D21089	Genbank	475156
Sequence 14304	D21235	Genbank	498145
Sequence 14305	D21262	Genbank	434764
Sequence 14306	D25215	Genbank	517114
Sequence 14307	D25274	Genbank	464185
Sequence 14308	D26600	Genbank	565650
Sequence 14309	D28468	Genbank	460704
J040000 1 1000			

Sequence 14310	D29011	Genbank	558525
Sequence 14311	D29643	Genbank	473936
Sequence 14312	D29805	Genbank	474986
Sequence 14313	D29954	Genbank	473940
Sequence 14314	D29958	Genbank	473948
Sequence 14315	D30648	Genbank	506337
•	D31885	Genbank	505097
Sequence 14316		Genbank	2366751
Sequence 14317	D32053	Genbank	644480
Sequence 14318	· D38112		633070
Sequence 14319	D38524	Genbank	
Sequence 14320	D42041	Genbank	577294
Sequence 14321	D42063	Genbank	924266
Sequence 14322	D49489	Genbank	1136742
Sequence 14323	D50420	Genbank	2618577
Sequence 14324	D50525	Genbank	1167502
Sequence 14325	D50916	Genbank	1469174
Sequence 14326	D50929	Genbank	1469200
Sequence 14327	D55673	Genbank	870746
Sequence 14328	D63486	Genbank	1469885
Sequence 14329	D63880	Genbank	961451
Sequence 14330	D64015	Genbank	2281005
Sequence 14331	D79205	Genbank	1754620
Sequence 14332	D79986	Genbank	1136389
Sequence 14333	D80000	Genbank	1136415
Sequence 14334	D83004	Genbank	1181557
Sequence 14335	D84273	Genbank	3894217
Sequence 14336	D86962	Genbank	1503997
Sequence 14337	D86985	Genbank	6634002
Sequence 14338	D87003	Genbank	2114242
Sequence 14339	D87127	Genbank	1817551
Sequence 14340	D87328	Genbank	1813423
	D87666	Genbank	1620016
Sequence 14341	D87684	Genbank	1663703
Sequence 14342		Genbank	220087
Sequence 14343	D90209	Genbank	2169143
Sequence 14344	E00882	Genbank	2169353
Sequence 14345	E01094	Genbank	2169333
Sequence 14346	E01533	G. G. 1.10 G. 1.11	
Sequence 14347	E01932	Genbank	2170180
Sequence 14348	E01954	Genbank	2170202
Sequence 14349	E01979	Genbank	2170227
Sequence 14350	E02164	Genbank	2170402
Sequence 14351	E02628	Genbank	2170856
Sequence 14352	E05647	Genbank	2173834
Sequence 14353	E05957	Genbank	2174144
Sequence 14354	E08663	Genbank	2176776
Sequence 14355	E12274	Genbank	3251108
Sequence 14356	E12457	Genbank	3251290
Sequence 14357	E12651	Genbank	3251483
Sequence 14358	E13330	Genbank	3252135
Sequence 14359	E13405	Genbank	3252210
Sequence 14360	J00139	Genbank	182715
Sequence 14361	J01415	Genbank	1944628
Sequence 14362	J02642	Genbank	182862
Sequence 14363	J02960	Genbank	178203
Sequence 14364	J03248	Genbank	183053
Sequence 14365	J03459	Genbank	187172
•			

Sequence 14366	J03537	Genbank	337513
Sequence 14367	J03592	Genbank	339722
Sequence 14368	J03746	Genbank	183655
Sequence 14369	J03779	Genbank	179833
Sequence 14370	J03934	Genbank	189245
Sequence 14371	J04173	Genbank	551173
Sequence 14372	J04801	Genbank	190371
Sequence 14373	J04991	Genbank	189425
Sequence 14374	J05633	Genbank	186504
Sequence 14375	K00558	Genbank	340020
Sequence 14376	K03432	Genbank	337377
Sequence 14377	L01042	Genbank	184097
Sequence 14378	L01057	Genbank	292462
Sequence 14379	L02320	Genbank	307365
Sequence 14380	L05094	Genbank	388033
Sequence 14381	L05491	Genbank	292831
Sequence 14382	L05628	Genbank	1835658
Sequence 14383	L06498	Genbank	292442
Sequence 14384	L07287	Genbank	292434
Sequence 14385	L07633	Genbank	186512
Sequence 14386	L08048	Genbank	184250
Sequence 14387	L08441	Genbank	179295
Sequence 14388	L09054	Genbank	307344
Sequence 14389	L10284	Genbank	186522
Sequence 14390	L10612	Genbank	402701
Sequence 14391	L11066	Genbank	307322
Sequence 14392	L11370	Genbank	387674
Sequence 14393	L12136	Genbank	181536
Sequence 14394	L12387	Genbank	459835
Sequence 14395	L12711	Genbank	388890
Sequence 14396	L13278	Genbank	292414
Sequence 14397	L14778	Genbank	306476
Sequence 14398	L19185	Genbank	440307
Sequence 14399	L19437	Genbank	4995970
Sequence 14400	L19739	Genbank	431318
Sequence 14401	L19783	Genbank	404725
Sequence 14402	L20046	Genbank	306741
Sequence 14403	L20941	Genbank	507251
Sequence 14404	L22569	Genbank	348706
Sequence 14405	L24444	Genbank	495860
Sequence 14406	L25931	Genbank	438638
Sequence 14407	L28809	Genbank	454151
Sequence 14408	L28997	Genbank	607027
Sequence 14409	L29766	Genbank	537525
Sequence 14410	L31610	Genbank	1220360
Sequence 14411	L37127	Genbank	4164098
Sequence 14412	L38995	Genbank	704415
Sequence 14413	L39320	Genbank	1226044
Sequence 14414	L39326	Genbank	1228094
Sequence 14415	L40396	Genbank	887357
Sequence 14416	L43575	Genbank	899064
Sequence 14417	L43964	Genbank	951202
Sequence 14418	L44140	Genbank	1203968
Sequence 14419	L47162	Genbank	1082035
Sequence 14420	L47647	Genbank	1000861
Sequence 14421	L76159	Genbank	1246232

Camuanaa 14400	L76416	Genbank	5566605
Sequence 14422	M10036	Genbank	339840
Sequence 14423		Genbank	182504
Sequence 14424	M11146	Genbank	182513
Sequence 14425	M11147		184092
Sequence 14426	M11353	Genbank	184233
Sequence 14427	M12623	Genbank	
Sequence 14428	M15205	Genbank	339718
Sequence 14429	M15796	Genbank	181271
Sequence 14430	M16541	Genbank	1280204
Sequence 14431	M17886	Genbank	190233
Sequence 14432	M18048	Genbank	337699
Sequence 14433	M19997	Genbank	181968
Sequence 14434	M21142	Genbank	183402
Sequence 14435	M21575	Genbank	1311702
Sequence 14436	M21895	Genbank	189523
Sequence 14437	M22760	Genbank	695359
Sequence 14438	M22918	Genbank	189019
Sequence 14439	M22920	Genbank	189021
Sequence 14440	M23410	Genbank	762884
Sequence 14441	M24070	Genbank	181485
Sequence 14442	M24194	Genbank	187701
Sequence 14443	M24486	Genbank	190785
Sequence 14444	M24543	Genbank	341200
Sequence 14445	M24690	Genbank	341218
Sequence 14446	M26481	Genbank	619789
Sequence 14447	M26663	Genbank	618463
Sequence 14448	M29064	Genbank	337452
	M29960	Genbank	339886
Sequence 14449	M31468	Genbank	190876
Sequence 14450	M31520	Genbank	337504
Sequence 14451		Genbank	178984
Sequence 14452	M36341	Genbank	189239
Sequence 14453	M36981	Genbank	179274
Sequence 14454	M37104	Genbank	184059
Sequence 14455	M37583	Genbank	189378
Sequence 14456	M38188		182392
Sequence 14457	M55150	Genbank	
Sequence 14458	M55409	Genbank	189596
Sequence 14459	M58485	Genbank	180154
Sequence 14460	M60457	Genbank	181249
Sequence 14461	M60484	Genbank	190225
Sequence 14462	M60756	Genbank	184085
Sequence 14463	M60857	Genbank	181334
Sequence 14464	M61764	Genbank	183702
Sequence 14465	M63180	Genbank	339679
Sequence 14466	M73547	Genbank	190161
Sequence 14467	M74775	Genbank	187151
Sequence 14468	M75099	Genbank	337369
Sequence 14469	M76299	Genbank	188778
Sequence 14470	M77232	Genbank	307392
Sequence 14471	M86181	Genbank	190566
Sequence 14472	M87339	Genbank	1498255
Sequence 14473	M88006	Genbank	178522
Sequence 14474	M88468	Genbank	307197
Sequence 14475	M90357	Genbank	179577
Sequence 14476	M93036	Genbank	182904
Sequence 14477	M93651	Genbank	338038

Sequence 14478	M94556	Genbank	188855
Sequence 14479	M96982	Genbank	338262
Sequence 14480	S48220	Genbank	257451
Sequence 14481	S54761	Genbank	265221
Sequence 14482	S72871	Genbank	639594
	S73498	Genbank	688010
Sequence 14483		Genbank	688296
Sequence 14484	S73591		
Sequence 14485	S74678	Genbank	241477
Sequence 14486	S81752	Genbank	1438780
Sequence 14487	S82597	Genbank	1699343
Sequence 14488	S82616	Genbank	1699307
Sequence 14489	U00238	Genbank	404860
Sequence 14490	U02031	Genbank	451329
Sequence 14491	U03886	Genbank	458225
Sequence 14492	U07231	Genbank	517195
Sequence 14493	U07857	Genbank	469048
Sequence 14494	U09466	Genbank	495492
	U09813	Genbank	1008454
Sequence 14495		Genbank	984280
Sequence 14496	U10248	=	
Sequence 14497	U10362	Genbank	505651
Sequence 14498	U12404	Genbank	531170
Sequence 14499	U12465	Genbank	562073
Sequence 14500	U12897	Genbank	562075
Sequence 14501	U13616	Genbank	608024
Sequence 14502	U14193	Genbank	555899
Sequence 14503	U14970	Genbank	550020
Sequence 14504	U14972	Genbank	550024
Sequence 14505	U14973	Genbank	550026
Sequence 14506	U17899	Genbank	717053
Sequence 14507	U18321	Genbank	603763
	U18671	Genbank	1293919
Sequence 14508		Genbank	644878
Sequence 14509	U20285		
Sequence 14510	U20362	Genbank	755485
Sequence 14511	U25182	Genbank	799380
Sequence 14512	U29171	Genbank	881618
Sequence 14513	U29589	Genbank	903978
Sequence 14514	U30826	Genbank	1049079
Sequence 14515	U34683	Genbank	1236349
Sequence 14516	U39817	Genbank	1072121
Sequence 14517	U41060	Genbank	1256000
Sequence 14518	U41195	Genbank	1244694
Sequence 14519	U41448	Genbank	3335023
Sequence 14520	U41668	Genbank	1477481
•	U44798	Genbank	1174216
Sequence 14521		Genbank	3077821
Sequence 14522	U46751	Genbank	
Sequence 14523	U49245	-	1216503
Sequence 14524	U51559	Genbank	1236709
Sequence 14525	U52111	Genbank	8331754
Sequence 14526	U52852	Genbank	8370402
Sequence 14527	U54993	Genbank	4097314
Sequence 14528	U56833	Genbank	1465750
Sequence 14529	U57846	Genbank	1373418
Sequence 14530	U57847	Genbank	1373420
Sequence 14531	U59911	Genbank	1403712
Sequence 14532	U66243	Genbank	1772645
Sequence 14533	U66589	Genbank	1575566
Coquence 17000	000000	Gonbank	

Sequence 14534	U67243	Genbank	2228516
Sequence 14535	U68566	Genbank	1916621
Sequence 14536	U68758	Genbank	4097815
Sequence 14537	U72648	Genbank	3914602
Sequence 14538	U73168	Genbank	1613899
Sequence 14539	U73645	Genbank	1737200
		Genbank	1857236
Sequence 14540	U73824		2315862
Sequence 14541	U75285	Genbank	
Sequence 14542	U76549	Genbank	1673574
Sequence 14543	U76669	Genbank	6672211
Sequence 14544	U77643	Genbank	2062390
Sequence 14545	U80017	Genbank	1737211
Sequence 14546	U80460	Genbank	2341006
Sequence 14547	U85625	Genbank	5091494
Sequence 14548	U87460	Genbank	2076881
Sequence 14549	U90904	Genbank	1913882
Sequence 14550	U90907	Genbank	1913886
Sequence 14551	U91327	Genbank	1871209
Sequence 14552	U91328	Genbank	2088550
Sequence 14553	U94586	Genbank	1946691
Sequence 14554	U95689	Genbank	2804669
Sequence 14555	U96629	Genbank	2341008
Sequence 14556	V00478	Genbank	28244
Sequence 14557	V00662	Genbank	13003
Sequence 14558	X00351	Genbank	28251
Sequence 14559	X02152	Genbank	34312
Sequence 14550	X03083	Genbank	34309
Sequence 14561	X03205	Genbank	36162
Sequence 14562	X05276	Genbank	37201
Sequence 14563	X05332	Genbank	35740
	X05857	Genbank	31988
Sequence 14564		Genbank	30204
Sequence 14565	X06409	Genbank	296365
Sequence 14566	X14608	Genbank	30146
Sequence 14567	X15822	Genbank	36129
Sequence 14568	X15940	Genbank	32355
Sequence 14569	X16135		
Sequence 14570	X16560	Genbank	30154
Sequence 14571	X17025	Genbank	488749
Sequence 14572	X17620	Genbank	35067
Sequence 14573	X52839	Genbank	36125
Sequence 14574	X52851	Genbank	30167
Sequence 14575	X52882	Genbank	311380
Sequence 14576	X52897	Genbank	24484
Sequence 14577	X52943	Genbank	28912
Sequence 14578	X56932	Genbank	23690
Sequence 14579	X56998	Genbank	37564
Sequence 14580	X58288	Genbank	32455
Sequence 14581	X59434	Genbank	432375
Sequence 14582	X60221	Genbank	509290
Sequence 14583	X61755	Genbank	32387
Sequence 14584	X63432	Genbank	28335
Sequence 14585	X63657	Genbank	296185
Sequence 14586	X64123	Genbank	35819
Sequence 14587	X65923	Genbank	31302
Sequence 14588	X66975	Genbank	32353
Sequence 14589	X67247	Genbank	36149
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<u>Table 3-5</u>

Sequence 14590	X67698	Genbank	37476
Sequence 14591	X69910	Genbank	297407
Sequence 14592	X70940	Genbank	38455
Sequence 14593	X71428	Genbank	393415
Sequence 14594	X72841	Genbank	297903
Sequence 14595	X74801	Genbank	671526
		Genbank	400462
Sequence 14596	X74979	<u> </u>	452319
Sequence 14597	X75593	Genbank	
Sequence 14598	X75684	Genbank	429092
Sequence 14599	X75861	Genbank	456258
Sequence 14600	X76013	Genbank	531595
Sequence 14601	X78137	Genbank	460770
Sequence 14602	X80200	Genbank	951276
Sequence 14603	X81713	Genbank	558213
Sequence 14604	X81788	Genbank	1045058
Sequence 14605	X81895	Genbank	2370158
Sequence 14606	X81900	Genbank	2274973
Sequence 14607	X84908	Genbank	1502344
Sequence 14608	X85372	Genbank	806563
Sequence 14609	X87241	Genbank	1107686
Sequence 14610	X87344	Genbank	1054740
Sequence 14611	X87949	Genbank	1143491
Sequence 14612	X91249	Genbank	1160185
Sequence 14613	X93036	Genbank	1085025
Sequence 14614	X94910	Genbank	3413292
Sequence 14615	X95592	Genbank	1185118
Sequence 14616	X96586	Genbank	1556398
Sequence 14617	X97544	Genbank	1770563
Sequence 14618	X98356	Genbank	1648869
	X99585	Genbank	1770518
Sequence 14619	Y00052	Genbank	30308
Sequence 14620		Genbank	36048
Sequence 14621	Y00282 Y00345	Genbank	35569
Sequence 14622		Genbank	34266
Sequence 14623	Y00815	Genbank	1552548
Sequence 14624	Y08201	Genbank	1743253
Sequence 14625	Y09836		2292905
Sequence 14626	Y10345	Genbank	
Sequence 14627	Y15286	Genbank	2584788
Sequence 14628	Z11692	Genbank	31107
Sequence 14629	Z11793	Genbank	36425
Sequence 14630	Z13009	Genbank	31072
Sequence 14631	Z14244	Genbank	30150
Sequence 14632	Z21507	Genbank	38521
Sequence 14633	Z24724	Genbank	505034
Sequence 14634	Z25749	Genbank	550116
Sequence 14635	Z31695	Genbank	469143
Sequence 14636	Z36832	Genbank	533946
Sequence 14637	Z36849	Genbank	533963
Sequence 14638	Z46973	Genbank	987947
Sequence 14639	Z47087	Genbank	860989
Sequence 14640	Z48950	Genbank	761715
Sequence 14641	Z54246	Genbank	1000700
Sequence 14642	Z68164	Genbank	1107692
Sequence 14643	Z68284	Genbank	1130698
Sequence 14644	Z69711	Genbank	1204109
Sequence 14645	Z69890	Genbank	1213361
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Sequence 14646	Z73900	Genbank	1340069
Sequence 14647	Z82242	Genbank	1841915
Sequence 14648	Z82975	Genbank	1673497
Sequence 14649	Z84474	Genbank	1903188
Sequence 14650	Z84812	Genbank	1845164
Sequence 14651	Z85986	Genbank	4034056
Sequence 14652	Z92544	Genbank	1869775
Sequence 14653	Z92845	Genbank	2462406
Sequence 14654	Z93041	Genbank	1905788
Sequence 14655	Z93928	Genbank	2578051
	Z95113	Genbank	6572188
Sequence 14656	Z95116	Genbank	2342581
Sequence 14657	Z95724	Genbank	2121307
Sequence 14658		Genbank	9650676
Sequence 14659	Z97053		2370077
Sequence 14660	Z97054	Genbank	2832593
Sequence 14661	Z97056	Genbank	
Sequence 14662	Z97353	Genbank	4455632
Sequence 14663	Z97652	Genbank	6456822
Sequence 14664	Z98049	Genbank	2462401
Sequence 14665	Z98749	Genbank	4775611
Sequence 14666	Z98884	Genbank	5304861
Sequence 14667	Z98950	Genbank	3036783
Sequence 14668	Z99128	Genbank	3135974
Sequence 14669	Z99916	Genbank	4158122
Sequence 14670	AA004759	EST	1448327
Sequence 14671	AA007319	EST	1463464
Sequence 14672	AA009421	EST	1470619
Sequence 14673	AA009598	EST	1470757
Sequence 14674	AA010071	EST	1471099
Sequence 14675	AA010798	EST	1471844
Sequence 14676	AA011369	EST	1472466
Sequence 14677	AA011539	EST	1472565
Sequence 14678	AA011569	EST	1472595
Sequence 14679	AA013065	EST	1474101
Sequence 14680	AA018720	EST	1481994
Sequence 14681	AA019036	EST	1482627
Sequence 14682	AA019504	EST	1482187
Sequence 14683	AA020851	EST	1484622
Sequence 14684	AA021306	EST	1484995
Sequence 14685	AA024597	EST	1489502
Sequence 14686	AA025057	EST	1489962
Sequence 14687	AA025037 AA025122	EST	1490037
Sequence 14688	AA026215	EST	1492901
	AA026316	EST	1492253
Sequence 14689	AA026316 AA026353	EST	1492674
Sequence 14690		EST	1492801
Sequence 14691	AA026643	EST	1492556
Sequence 14692	AA026758		1492149
Sequence 14693	AA027229	EST	
Sequence 14694	AA029679	EST	1497082
Sequence 14695	AA034414	EST	1506241
Sequence 14696	AA035260	EST	1506772
Sequence 14697	AA035377	EST	1507101
Sequence 14698	AA037143	EST	1512251
Sequence 14699	AA039589	EST	1516014
Sequence 14700	AA039683	EST	1516155
Sequence 14701	AA039734	EST	1516236
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Sequence 14702	AA039876	EST	1516172
Sequence 14703	AA039929	EST	1516206
Sequence 14704	AA040404	EST	1516765
Sequence 14705	AA043048	EST	1522583
Sequence 14706	AA043193	EST	1521111
Sequence 14707	AA043606	EST	1521472
Sequence 14708	AA044663	EST	1523043
	AA044835	EST	1523038
Sequence 14709	AA046542	EST	1524439
Sequence 14710		EST	1524915
Sequence 14711	AA046810	EST	1524763
Sequence 14712	AA046864		1524944
Sequence 14713	AA047046	EST	1544534
Sequence 14714	AA053607	EST	
Sequence 14715	AA053853	EST	1544797
Sequence 14716	AA053926	EST	1544870
Sequence 14717	AA053942	EST	1545004
Sequence 14718	AA054471	EST	1545396
Sequence 14719	AA056107	EST	1548444
Sequence 14720	AA057352	EST	1549991
Sequence 14721	AA057541	EST	1550246
Sequence 14722	AA057559	EST	1550199
Sequence 14723	AA063079	EST	1556632
Sequence 14724	AA063612	EST	1557579
Sequence 14725	AA065328	EST	1929208
Sequence 14726	AA069023	EST	1576401
Sequence 14727	AA069422	EST	1576781
Sequence 14728	AA069655	EST	1577014
Sequence 14729	AA070277	EST	1577637
Sequence 14730	AA070855	EST	1578487
Sequence 14731	AA071269	EST	1578757
Sequence 14732	AA071402	EST	1578773
Sequence 14733	AA071472	EST	1578903
Sequence 14734	AA074221	EST	1614090
Sequence 14735	AA074611	EST	1614677
Sequence 14736	AA074614	EST	1614492
Sequence 14737	AA074664	EST	1614608
Sequence 14737 Sequence 14738	AA074831	EST	1614875
Sequence 14739	AA075200	EST	1615070
	AA075200	EST	1615850
Sequence 14740	AA079050	EST	1617942
Sequence 14741	AA081478	EST	1623536
Sequence 14742	AA081476 AA082545	EST	1624602
Sequence 14743	AA082343 AA083712	EST	1625772
Sequence 14744		EST	1626735
Sequence 14745	AA084678	EST	1626988
Sequence 14746	AA084868	EST	1633764
Sequence 14747	AA088189		1636589
Sequence 14748	AA090105	EST EST	1637531
Sequence 14749	AA092310		1637545
Sequence 14750	AA092324	EST	1644862
Sequence 14751	AA098978	EST	1644895
Sequence 14752	AA099015	EST	
Sequence 14753	AA099080	EST	1645522
Sequence 14754	AA099327	EST	1646033
Sequence 14755	AA100639	EST	1646940
Sequence 14756	AA101116	EST	1647692
Sequence 14757	AA101598	EST	1648552

Sequence 14758	AA101663	EST	1648744
Sequence 14759	AA102442	EST	1647356
Sequence 14760	AA102560	EST	1647752
Sequence 14761	AA114042	EST	1667919
Sequence 14762	AA115474	EST	1670145
Sequence 14763	AA126478	EST	1686071
Sequence 14764	AA126762	EST	1686262
Sequence 14765	AA126841	EST	1686323
Sequence 14766	AA127483	EST	1686755
Sequence 14767	AA127550	EST	1686822
Sequence 14768	AA127886	EST	1687193
Sequence 14769	AA129205	EST	1688990
Sequence 14770	AA129465	EST	1689336
Sequence 14770	AA129935	EST	1689511
	AA131336	EST	1692834
Sequence 14772	AA132089	EST	1693579
Sequence 14773		EST	1693885
Sequence 14774	AA132394	EST	1694505
Sequence 14775	AA132989		1694539
Sequence 14776	AA133031	EST	1690768
Sequence 14777	AA133736	EST	
Sequence 14778	AA133759	EST	1690727
Sequence 14779	AA134263	EST	1691637
Sequence 14780	AA134609	EST	1695597
Sequence 14781	AA135030	EST	1696141
Sequence 14782	AA135126	EST	1696227
Sequence 14783	AA135254	EST	1696373
Sequence 14784	AA136337	EST	1697545
Sequence 14785	AA136723	EST	1697933
Sequence 14786	AA136761	EST	1698008
Sequence 14787	AA137016	EST	1698225
Sequence 14788	AA142942	EST	1712320
Sequence 14789	AA143324	EST	1712695
Sequence 14790	AA143422	EST	1712852
Sequence 14791	AA147059	EST	1716433
Sequence 14792	AA147948	EST	1717455
Sequence 14793	AA148889	EST	1718998
Sequence 14794	AA149404	EST	1719920
Sequence 14795	AA149518	EST	1720336
Sequence 14796	AA149579	EST	1720380
Sequence 14797	AA149801	EST	1720899
Sequence 14798	AA149826	EST	1720961
Sequence 14799	AA150361	EST	1721873
Sequence 14800	AA150481	EST	1721995
Sequence 14801	AA151140	EST	1719383
Sequence 14802	AA151423	EST	1719820
Sequence 14803	AA151795	EST	1720490
Sequence 14804	AA151809	EST	1720504
Sequence 14805	AA151816	EST	1720511
Sequence 14806	AA152179	EST	1721231
Sequence 14807	AA155999	EST	1727666
Sequence 14808	AA156226	EST	1727844
Sequence 14809	AA158153	EST	1732990
Sequence 14810	AA158669	EST	1733480
Sequence 14811	AA158704	EST	1733533
Sequence 14812	AA159176	EST	1733969
Sequence 14813	AA159347	EST	1733909
Sequence 14013	AA 108041	LOI	170-71-73

Sequence 14814	AA160666	EST	1736043
Sequence 14815	AA161467	EST	1735906
Sequence 14816	AA165437	EST	1741453
Sequence 14817	AA166691	EST	1745155
Sequence 14818	AA166843	EST	1745024
	AA166946	EST	1745322
Sequence 14819		EST	1745386
Sequence 14820	AA167011		
Sequence 14821	AA167501	EST	1745886
Sequence 14822	AA169786	EST	1748121
Sequence 14823	AA171706	EST	1750791
Sequence 14824	AA173421	EST	1753550
Sequence 14825	AA173705	EST	1753883
Sequence 14826	AA173936	EST	1754068
Sequence 14827	AA176244	EST	1757375
Sequence 14828	AA176455	EST	1757713
Sequence 14829	AA178955	EST	1760308
Sequence 14830	AA179220	EST	1760572
Sequence 14831	AA181891	EST	1765368
Sequence 14832	AA182751	EST	1766581
Sequence 14833	AA182940	EST	1766108
	AA186328	EST	1774446
Sequence 14834		EST	1775103
Sequence 14835	AA186811	EST	1779239
Sequence 14836	AA190408		
Sequence 14837	AA192099	EST	1781923
Sequence 14838	AA192347	EST	1781585
Sequence 14839	AA194517	EST	1784213
Sequence 14840	AA195093	EST	1784783
Sequence 14841	AA203477	EST	1799204
Sequence 14842	AA204856	EST	1802881
Sequence 14843	AA205970	EST	1801341
Sequence 14844	AA206446	EST	1801826
Sequence 14845	AA206761	EST	1802148
Sequence 14846	AA210909	EST	1809555
Sequence 14847	AA214548	EST	1813173
Sequence 14848	AA215640	EST	1815385
Sequence 14849	AA216047	EST	1816018
Sequence 14850		EST	1816048
Sequence 14851	AA219137	EST	1833202
Sequence 14852	AA220941	EST	1839687
Sequence 14853	AA223990	EST	1844515
•		EST	1846276
Sequence 14854		EST	1846765
Sequence 14855		EST	1846812
Sequence 14856			1846970
Sequence 14857	AA225653	EST	
Sequence 14858		EST	1847200
Sequence 14859		EST	1848466
Sequence 14860		EST	1848839
Sequence 14861	AA229713	EST	1851876
Sequence 14862		EST	1855538
Sequence 14863		EST	1860273
Sequence 14864		EST	1874187
Sequence 14865		EST	1874456
Sequence 14866		EST	1874594
Sequence 14867		EST	1875185
Sequence 14868		EST	1879338
Sequence 14869		EST	1886090
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Sequence 14870	AA252998	EST	1882921
Sequence 14871	AA257006	EST	1891135
Sequence 14872	AA258030	EST	1894480
Sequence 14873	AA258271	EST	1893453
Sequence 14874	AA258394	EST	1893536
	AA261894	EST	1897728
Sequence 14875			
Sequence 14876	AA261960	EST	1898149
Sequence 14877	AA262181	EST	1898452
Sequence 14878	AA262425	EST	1897921
Sequence 14879	AA262871	EST	1898592
Sequence 14880	AA263047	EST	1898845
Sequence 14881	AA278243	EST	1920183
Sequence 14882	AA278787	EST	1920108
Sequence 14883	AA279139	EST	1920605
Sequence 14884	AA279421	EST	1920886
Sequence 14885	AA280206	EST	1921881
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Sequence 11677: Found in patent publication WO99/46375

Sequence 11679: Found in patent publication WO99/63088

Sequence 11680: Found in patent publication WO99/67383

Sequence 13322: Found in patent publication WO00/09552

Sequence 13323: Found in patent publication WO99/66041

Sequence 13324: Found in patent publication WO99/66041

Sequence 13325: Found in patent publication WO99/67383

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Sequence 13326: Found in patent publication WO00/06698

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Sequence 13327: Found in patent publication US6034229

Sequence 13328: Found in patent publication WO00/18925

Sequence 15908: Found in patent publication WO98/33916

Sequence 15909: Found in patent publication WO99/14328

Sequence 15910: Found in patent publication WO99/66041

Sequence 15911: Found in patent publication WO00/05382

Sequence 11676: Found in patent publication WO00/05367

Sequence 11677: Found in patent publication WO99/46375

Sequence 11679: Found in patent publication WO99/63088

Sequence 11680: Found in patent publication WO99/67383

Sequence 13322: Found in patent publication WO00/09552

Sequence 13323: Found in patent publication WO99/66041

Sequence 13324: Found in patent publication WO99/66041

Sequence 13325: Found in patent publication WO99/67383

AGTACTTCTAGAATTAATTAAGGCGGGTCCCTGCTTGTCCTATGAACTGCTCAGAGCTCC
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Sequence 13326: Found in patent publication WO00/06698

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Sequence 13327: Found in patent publication US6034229

Sequence 13328: Found in patent publication WO00/18925

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Sequence 15908: Found in patent publication WO98/33916

Sequence 15909: Found in patent publication WO99/14328

Sequence 15910: Found in patent publication WO99/66041

Sequence 15911: Found in patent publication WO00/05382

Sequence 11676: Found in patent publication WO00/05367

Sequence 11677: Found in patent publication WO99/46375

Sequence 11679: Found in patent publication WO99/63088

Sequence 11680: Found in patent publication WO99/67383

Sequence 13322: Found in patent publication WO00/09552

Sequence 13323: Found in patent publication WO99/66041

Sequence 13324: Found in patent publication WO99/66041

Sequence 13325: Found in patent publication WO99/67383

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Sequence 13326: Found in patent publication WO00/06698

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Sequence 15908: Found in patent publication WO98/33916

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Sequence 15910: Found in patent publication WO99/66041

Sequence 15911: Found in patent publication WO00/05382

Sequence 11676: Found in patent publication WO00/05367

Sequence 11677: Found in patent publication WO99/46375

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Sequence 13325: Found in patent publication WO99/67383

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Sequence 13326: Found in patent publication WO00/06698

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Sequence 13324: Found in patent publication WO99/66041

Sequence 13325: Found in patent publication WO99/67383

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Sequence 13326: Found in patent publication WO00/06698

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Sequence 13327: Found in patent publication US6034229

Sequence 13328: Found in patent publication WO00/18925

Sequence 15908: Found in patent publication WO98/33916

Sequence 15909: Found in patent publication WO99/14328

Sequence 15910: Found in patent publication WO99/66041

Sequence 15911: Found in patent publication WO00/05382

Sequence 11676: Found in patent publication WO00/05367

Sequence 11677: Found in patent publication WO99/46375

Sequence 11679: Found in patent publication WO99/63088

Sequence 11680: Found in patent publication WO99/67383

Sequence 13322: Found in patent publication WO00/09552

Sequence 13323: Found in patent publication WO99/66041

Sequence 13324: Found in patent publication WO99/66041

Sequence 13325: Found in patent publication WO99/67383

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Table 4-1

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Sequence 5431	M58581	Sequence 5483	U33821
Sequence 5432	M59465	Sequence 5484	U34252
Sequence 5433	M62401	Sequence 5485	U37230
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Sequence 5435	M65131	Sequence 5487	U39400
Sequence 5436	M69226	Sequence 5488	U40272
Sequence 5437	M69238	Sequence 5489	U42404
Sequence 5438	M74091	Sequence 5490	U43899
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Sequence 5442	M82919	Sequence 5494	U57057
Sequence 5443	M84443	Sequence 5495	U58855
Sequence 5444	M86917	Sequence 5496	U59919
Sequence 5445	M90054	Sequence 5497	U60205
Sequence 5446	M90309	Sequence 5498	U61232
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Sequence 5450	M96803	Sequence 5502	U68758
Sequence 5451	S50223	Sequence 5503	U70063
Sequence 5452	S57235	Sequence 5504	U70322
Sequence 5453	S59049	Sequence 5505	U70451
Sequence 5454	S64650	Sequence 5506	U72761
Sequence 5455	S68015	Sequence 5507	U75272
Sequence 5456	S69272	Sequence 5508	U75679
Sequence 5457	S73498	Sequence 5509	U78575
Sequence 5458	U02680	Sequence 5510	U80034
Sequence 5459	U05291	Sequence 5511	U83115

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Sequence 5513	U87166	Sequence 5565	X97675
Sequence 5514	U90915	Sequence 5566	X98263
Sequence 5515	U96759	Sequence 5567	Y00282
Sequence 5516	V00478	Sequence 5568	Y00345
Sequence 5517	X02530	Sequence 5569	Y08991
Sequence 5518	X04098	Sequence 5570	Y09615
Sequence 5519	X05332	Sequence 5571	Y09836
Sequence 5520	X05345	Sequence 5572	Y12065
Sequence 5521	X06272	Sequence 5573	Y12860
Sequence 5522	X06617	Sequence 5574	Y13286
Sequence 5523	X12433	Sequence 5575	Y15286
Sequence 5524	X12791	Sequence 5576	Y16132
Sequence 5525	X13425	Sequence 5577	Y16241
Sequence 5526	X13482	Sequence 5578	Y16414
Sequence 5527	X13585	Sequence 5579	Y18314
Sequence 5528	X13839	Sequence 5580	Z21507
Sequence 5529	X15187	Sequence 5581	Z22658
Sequence 5530	X15729	Sequence 5582	Z36531
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Sequence 5533	X53793	Sequence 5585	Z47553
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Sequence 5538	X62534	Sequence 5590	T84934
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Sequence 5548	X76771	Sequence 5600	X26850
Sequence 5549	X77196	Sequence 5601	X27262
Sequence 5550	X78627	Sequence 5602	X29296
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Sequence 5560	X87176	Sequence 5612	Z14392
Sequence 5561	X87949	Sequence 5613	Z15132
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Sequence 5563	X95384	Sequence 5615	A06919

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Sequence 5618	AB002375	Sequence 5670	AL049932
Sequence 5619	AB006534	Sequence 5671	AL117499
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Sequence 5621	AB007899	Sequence 5673	D38551
Sequence 5622	AB011079	Sequence 5674	D42044
Sequence 5623	AB014458	Sequence 5675	D49396
Sequence 5624	AB015639	Sequence 5676	D50371
Sequence 5625	AB016068	Sequence 5677	D50525
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Sequence 5634	AB033114	Sequence 5686	L15203
Sequence 5635	AF007791	Sequence 5687	L34839
Sequence 5636	AF015040	Sequence 5688	M10119
Sequence 5637	AF016371	Sequence 5689	M13231
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Sequence 5642	AF052642	Sequence 5694	M24902
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Sequence 5644	AF067008	Sequence 5696	M74777
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Sequence 5646	AF070561	Sequence 5698	S69272
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Sequence 5648	AF077202	Sequence 5700	S75755
Sequence 5649	AF078854	Sequence 5701	U07919
Sequence 5650	AF085845	Sequence 5702	U21936
Sequence 5651	AF093097	Sequence 5703	U40272
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Sequence 5653	AF100759	Sequence 5705	U79751
Sequence 5654	AF103907	Sequence 5706	U96759
Sequence 5655	AF104914	Sequence 5707	X05606
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Sequence 5664	AF159056	Sequence 5716	Y00345
Sequence 5665	AF188746	Sequence 5717	Z11793
Sequence 5666	AF191298	Sequence 5718	AA026715
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Sequence 5726	AA100486	Sequence 5778	AA506953
Sequence 5727	AA128739	Sequence 5779	AA507804
Sequence 5728	AA133307	Sequence 5780	AA515143
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Sequence 5730	AA148729	Sequence 5782	AA541587
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Sequence 5746	AA262783	Sequence 5798	AA779728
Sequence 5747	AA278475	Sequence 5799	AA806808
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Sequence 5760	AA393313	Sequence 5812	AI049995
Sequence 5761	AA400692	Sequence 5813	AI084651
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Sequence 5766	AA454557	Sequence 5818	Al133727
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Sequence 5768	AA468839	Sequence 5820	Al282318
Sequence 5769	AA478518	Sequence 5821	Al300489
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Sequence 5839	AL037316	Sequence 5891	AA011070
Sequence 5840	AL037732	Sequence 5892	AA013090
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Sequence 5847	AW173228	Sequence 5899	AA024796
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Sequence 5852	R24400	Sequence 5904	AA029070
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Sequence 5855	R78584	Sequence 5907	AA029676
Sequence 5856	T81995	Sequence 5908	AA032220
Sequence 5857	T99386	Sequence 5909	AA033651
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Sequence 5859	T59274	Sequence 5911	AA034979
Sequence 5860	X06834	Sequence 5912	AA035790
Sequence 5861	X24060	Sequence 5913	AA037210
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Sequence 5863	X52268	Sequence 5915	AA037886
Sequence 5864	Z13254	Sequence 5916	AA039381
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Sequence 5866	AC003991	Sequence 5918	AA043137
Sequence 5867	AC007275	Sequence 5919	AA043159
Sequence 5868	AP000502	Sequence 5920	AA043415
Sequence 5869	AC007239	Sequence 5921	AA044239
Sequence 5870	AC000065	Sequence 5922	AA044941
Sequence 5871	AC005881	Sequence 5923	AA045147
Sequence 5872	AC004982	Sequence 5924	AA045410
Sequence 5873	AC003991	Sequence 5925	AA045803
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Sequence 5930	AA054492	Sequence 5982	AA100852
Sequence 5931	AA056060	Sequence 5983	AA101777
Sequence 5932	AA056358	Sequence 5984	AA112626
Sequence 5933	AA056365	Sequence 5985	AA113403
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Sequence 5941	AA062583	Sequence 5993	AA128114
Sequence 5942	AA062994	Sequence 5994	AA128647
Sequence 5943	AA063247	Sequence 5995	AA128668
Sequence 5944	AA069367	Sequence 5996	AA129772
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Sequence 5949	AA074445	Sequence 6001	AA131252
Sequence 5950	AA074509	Sequence 6002	AA131284
Sequence 5951	AA075460	Sequence 6003	AA131793
Sequence 5952	AA076295	Sequence 6004	AA132510
Sequence 5953	AA081007	Sequence 6005	AA132792
Sequence 5954	AA081355	Sequence 6006	AA133068
Sequence 5955	AA082209	Sequence 6007	AA133268
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Sequence 5957	AA083410	Sequence 6009	AA134701
Sequence 5958	AA083411	Sequence 6010	AA135870
Sequence 5959	AA083482	Sequence 6011	AA137165
Sequence 5960	AA083510	Sequence 6012	AA143242
Sequence 5961	AA083772	Sequence 6013	AA143523
Sequence 5962	AA083836	Sequence 6014	AA143548
Sequence 5963	AA084221	Sequence 6015	AA146582
Sequence 5964	AA084698	Sequence 6016	AA146900
Sequence 5965	AA085480	Sequence 6017	AA147608
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Sequence 5967	AA088679	Sequence 6019	AA148019
Sequence 5968	AA088758	Sequence 6020	AA148269
Sequence 5969	AA090106	Sequence 6021	AA148569
Sequence 5970	AA090203	Sequence 6022	AA149490
Sequence 5971	AA090548	Sequence 6023	AA149543
Sequence 5972	AA091275	Sequence 6024	AA149551
Sequence 5973	AA092060	Sequence 6025	AA149579
Sequence 5974	AA092343	Sequence 6026	AA149852
Sequence 5975	AA092532	Sequence 6027	AA150480
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Sequence 5977	AA093935	Sequence 6029	AA150780
Sequence 5978	AA094331	Sequence 6030	AA150891
Sequence 5979	AA094656	Sequence 6031	AA151596

Table 4-1

Sequence 6032	AA152073	Sequence 6084	AA203206
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Sequence 6035	AA155828	Sequence 6087	AA205033
Sequence 6036	AA155865	Sequence 6088	AA205647
Sequence 6037	AA156369	Sequence 6089	AA205857
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Sequence 6039	AA159272	Sequence 6091	AA210760
Sequence 6040	AA159647	Sequence 6092	AA214668
Sequence 6041	AA159771	Sequence 6093	AA214688
Sequence 6042	AA160677	Sequence 6094	AA218788
Sequence 6043	AA165027	Sequence 6095	AA223249
Sequence 6044	AA165082	Sequence 6096	AA223433
Sequence 6045	AA165165	Sequence 6097	AA223956
Sequence 6046	AA165632	Sequence 6098	AA224124
Sequence 6047	AA166707	Sequence 6099	AA224994
Sequence 6048	AA167399	Sequence 6100	AA225115
Sequence 6049	AA169507	Sequence 6101	AA225294
Sequence 6050	AA171510	Sequence 6102	AA225488
Sequence 6051	AA173391	Sequence 6103	AA226171
Sequence 6052	AA173559	Sequence 6104	AA226234
Sequence 6053	AA177050	Sequence 6105	AA226321
Sequence 6054	AA179187	Sequence 6106	AA228273
Sequence 6055	AA179462	Sequence 6107	AA228431
Sequence 6056	AA179487	Sequence 6108	AA228822
Sequence 6057	AA180137	Sequence 6109	AA228940
Sequence 6058	AA180154	Sequence 6110	AA229495
Sequence 6059	AA181618	Sequence 6111	AA229611
Sequence 6060	AA182457	Sequence 6112	AA232691
Sequence 6061	AA186538	Sequence 6113	AA232700
Sequence 6062	AA187928	Sequence 6114	AA233073
Sequence 6063	AA187998	Sequence 6115	AA233088
Sequence 6064	AA188052	Sequence 6116	AA233131
Sequence 6065	AA188312	Sequence 6117	AA234073
Sequence 6066	AA188396	Sequence 6118	AA234092
Sequençe 6067	AA188875	Sequence 6119	AA234990
Sequence 6068	AA189150	Sequence 6120	AA235914
Sequence 6069	AA190529	Sequence 6121	AA236320
Sequence 6070	AA190661	Sequence 6122	AA242891
Sequence 6071	AA190843	Sequence 6123	AA243143
Sequence 6072	AA190890	Sequence 6124	AA243696
Sequence 6073	AA191045	Sequence 6125	AA244003
Sequence 6074	AA191719	Sequence 6126	AA244116
Sequence 6075	AA191721	Sequence 6127	AA244207
Sequence 6076	AA192597	Sequence 6128	AA250748
Sequence 6077	AA194062	Sequence 6129	AA251178
Sequence 6078	AA194988	Sequence 6130	AA251578
Sequence 6079	AA196001	Sequence 6131	AA251644
Sequence 6080	AA196392	Sequence 6132	AA252849
Sequence 6081	AA196978	Sequence 6133	AA253453
Sequence 6082	AA199658	Sequence 6134	AA255761
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Table 4-1

Sequence 6136	AA257973	Sequence 6188	AA312059
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Sequence 6139	AA261990	Sequence 6191	AA313057
Sequence 6140	AA262513	Sequence 6192	AA313244
Sequence 6141	AA263153	Sequence 6193	AA313653
Sequence 6142	AA278520	Sequence 6194	AA313979
Sequence 6143	AA279517	Sequence 6195	AA314659
Sequence 6144	AA280917	Sequence 6196	AA314810
Sequence 6145	AA284615	Sequence 6197	AA315016
Sequence 6146	AA286953	Sequence 6198	AA316411
Sequence 6147	AA287511	Sequence 6199	AA317068
Sequence 6148	AA291325	Sequence 6200	AA317496
Sequence 6149	AA291363	Sequence 6201	AA317976
Sequence 6150	AA291602	Sequence 6202	AA320004
Sequence 6151	AA291764	Sequence 6203	AA320723
Sequence 6152	AA292638	Sequence 6204	AA322540
Sequence 6153	AA293028	Sequence 6205	AA324478
Sequence 6154	AA293629	Sequence 6206	AA326306
Sequence 6155	AA295182	Sequence 6207	AA327105
Sequence 6156	AA295788	Sequence 6208	AA327469
Sequence 6157	AA296846	Sequence 6209	AA328770
Sequence 6158	AA297941	Sequence 6210	AA329242
Sequence 6159	AA298085	Sequence 6211	AA329770
Sequence 6160	AA298104	Sequence 6212	AA329937
Sequence 6161	AA298294	Sequence 6213	AA330123
Sequence 6162	AA298464	Sequence 6214	AA332083
Sequence 6163	AA298489	Sequence 6215	AA332593
Sequence 6164	AA300065	Sequence 6216	AA332716
Sequence 6165	AA302457	Sequence 6217	AA333090
Sequence 6166	AA305358	Sequence 6218	AA333526
Sequence 6167	AA305627	Sequence 6219	AA333765
Sequence 6168	AA305824	Sequence 6220	AA334344
Sequence 6169	AA305895	Sequence 6221	AA338215
Sequence 6170	AA305909	Sequence 6222	AA340635
Sequence 6171	AA306264	Sequence 6223	AA344625
Sequence 6172	AA306413	Sequence 6224	AA349232
Sequence 6173	AA306480	Sequence 6225	AA351297
Sequence 6174	AA306573	Sequence 6226	AA352938
Sequence 6175	AA306812	Sequence 6227	AA354872
Sequence 6176	AA306956	Sequence 6228	AA358790
Sequence 6177	AA306982	Sequence 6229	AA359838
Sequence 6178	AA306983	Sequence 6230	AA360190
Sequence 6179	AA307513	Sequence 6231	AA360310
Sequence 6180	AA307697	Sequence 6232	AA364171
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Sequence 6182	AA309909	Sequence 6234	AA365620
Sequence 6183	AA310173	Sequence 6235	AA366995
Sequence 6184	AA310807	Sequence 6236	AA367471
Sequence 6185	AA311028	Sequence 6237	AA370110
Sequence 6186	AA311905	Sequence 6238	AA370222
Sequence 6187	AA311978	Sequence 6239	AA370352

Sequence 6240	AA370507	Sequence 6292	AA442976
Sequence 6241	AA370620	Sequence 6293	AA443024
Sequence 6242	AA371017	Sequence 6294	AA443082
Sequence 6243	AA371157	Sequence 6295	AA443242
Sequence 6244	AA371265	Sequence 6296	AA444093
Sequence 6245	AA374020	Sequence 6297	AA446298
Sequence 6246	AA374198	Sequence 6298	AA446910
Sequence 6247	AA374814	Sequence 6299	AA447608
Sequence 6248	AA375927	Sequence 6300	AA447948
Sequence 6249	AA376034	Sequence 6301	AA448795
Sequence 6250	AA378933	Sequence 6302	AA448882
Sequence 6251	AA379448	Sequence 6303	AA449520
Sequence 6252	AA397832	Sequence 6304	AA452012
Sequence 6253	AA397953	Sequence 6305	AA452027
Sequence 6254	AA398732	Sequence 6306	AA452352
Sequence 6255	AA399275	Sequence 6307	AA453352
Sequence 6256	AA400696	Sequence 6308	AA453651
Sequence 6257	AA401206	Sequence 6309	AA453719
Sequence 6258	AA402115	Sequence 6310	AA453784
Sequence 6259	AA403120	Sequence 6311	AA454187
Sequence 6260	AA403121	Sequence 6312	AA454962
Sequence 6261	AA403188	Sequence 6313	AA455668
Sequence 6262	AA404444	Sequence 6314	AA455894
Sequence 6263	AA416736	Sequence 6315	AA456390
Sequence 6264	AA417876	Sequence 6316	AA458761
Sequence 6265	AA418628	Sequence 6317	AA458770
Sequence 6266	AA419263	Sequence 6318	AA459167
Sequence 6267	AA419554	Sequence 6319	AA459210
Sequence 6268	AA420571	Sequence 6320	AA461128
Sequence 6269	AA420633	Sequence 6321	AA461467
Sequence 6270	AA420705	Sequence 6322	AA461605
Sequence 6271	AA421682	Sequence 6323	AA464133
Sequence 6272	AA421850	Sequence 6324	AA464141
Sequence 6273	AA422060	Sequence 6325	AA467750
Sequence 6274	AA423801	Sequence 6326	AA468167
Sequence 6275	AA425407	Sequence 6327	AA468404
Sequence 6276	AA425526	Sequence 6328	AA468624
Sequence 6277	AA426578	Sequence 6329	AA468839
Sequence 6278	AA427866	Sequence 6330	AA469129
Sequence 6279	AA428008	Sequence 6331	AA469303
Sequence 6280	AA428084	Sequence 6332	AA469345
Sequence 6281	AA429014	Sequence 6333	AA469406
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Sequence 6283	AA429493	Sequence 6335	AA476758
Sequence 6284	AA430110	Sequence 6336	AA479167
Sequence 6285	AA431677	Sequence 6337	AA479280
Sequence 6286	AA432103	Sequence 6338	AA479287
Sequence 6287	AA433869	Sequence 6339	AA479646
Sequence 6288	AA433988	Sequence 6340	AA481078
Sequence 6289	AA435565	Sequence 6341	AA481316
Sequence 6290	AA437224	Sequence 6342	AA482539
Sequence 6291	AA442184	Sequence 6343	AA482724

Table 4-1

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Sequence 6346	AA487448	Sequence 6398	AA528202
Sequence 6347	AA487747	Sequence 6399	AA531221
Sequence 6348	AA488633	Sequence 6400	AA531312
Sequence 6349	AA492035	Sequence 6401	AA531509
Sequence 6350	AA492280	Sequence 6402	AA531563
Sequence 6351	AA493222	Sequence 6403	AA531602
Sequence 6352	AA493331	Sequence 6404	AA533057
Sequence 6353	AA493448	Sequence 6405	AA533648
Sequence 6354	AA493512	Sequence 6406	AA534235
Sequence 6355	AA493956	Sequence 6407	AA534281
Sequence 6356	AA494214	Sequence 6408	AA534435
Sequence 6357	AA494493	Sequence 6409	AA535377
Sequence 6358	AA495759	Sequence 6410	AA536036
Sequence 6359	AA496957	Sequence 6411	AA542981
Sequence 6360	AA501363	Sequence 6412	AA548472
Sequence 6361	AA501968	Sequence 6413	AA548722
Sequence 6362	AA502901	Sequence 6414	AA551464
Sequence 6363	AA502979	Sequence 6415	AA552715
Sequence 6364	AA503020	Sequence 6416	AA554661
Sequence 6365	AA504334	Sequence 6417	AA554735
Sequence 6366	AA504953	Sequence 6418	AA557888
Sequence 6367	AA505523	Sequence 6419	AA559906
Sequence 6368	AA505953	Sequence 6420	AA564484
Sequence 6369	AA507442	Sequence 6421	AA564505
Sequence 6370	AA507777	Sequence 6422	AA573742
Sequence 6371	AA508889	Sequence 6423	AA573762
Sequence 6372	AA512974	Sequence 6424	AA575882
Sequence 6373	AA514938	Sequence 6425	AA576425
Sequence 6374	AA515132	Sequence 6426	AA577553
Sequence 6375	AA515143	Sequence 6427	AA578209
Sequence 6376	AA515564	Sequence 6428	AA578511
Sequence 6377	AA516531	Sequence 6429	AA578773
Sequence 6378	AA522936	Sequence 6430	AA578881
Sequence 6379	AA523252	Sequence 6431	AA578976
Sequence 6380	AA523464	Sequence 6432	AA579384
Sequence 6381	AA523498	Sequence 6433	AA579486
Sequence 6382	AA523652	Sequence 6434	AA582591
Sequence 6383	AA523902	Sequence 6435	AA583567
Sequence 6384	AA524485	Sequence 6436	AA583574
Sequence 6385	AA524778	Sequence 6437	AA587140
Sequence 6386	AA525419	Sequence 6438	AA587226
Sequence 6387	AA526515	Sequence 6439	AA587630
Sequence 6388	AA526591	Sequence 6440	AA592908
Sequence 6389	AA526886	Sequence 6441	AA594681
Sequence 6390	AA527187	Sequence 6442	AA594682
Sequence 6391	AA527272	Sequence 6443	AA595471
Sequence 6392	AA527730	Sequence 6444	AA602794
Sequence 6393	AA527737	Sequence 6445	AA602957
Sequence 6394	AA527805	Sequence 6446	AA603041
Sequence 6395	AA528023	Sequence 6447	AA603135

Sequence 6448	AA604474	Sequence 6500	AA699999
Sequence 6449	AA604497	Sequence 6501	AA702398
Sequence 6450	AA605077	Sequence 6502	AA702888
Sequence 6451	AA609094	Sequence 6503	AA705447
Sequence 6452	AA609436	Sequence 6504	AA705736
Sequence 6453	AA610476	Sequence 6505	AA705743
Sequence 6454	AA613916	Sequence 6506	AA705813
Sequence 6455	AA613933	Sequence 6507	AA706241
Sequence 6456	AA614105	Sequence 6508	AA707490
Sequence 6457	AA614525	Sequence 6509	AA708143
Sequence 6458	AA618484	Sequence 6510	AA708195
Sequence 6459	AA620750	Sequence 6511	AA713687
Sequence 6460	AA620995	Sequence 6512	AA714934
Sequence 6461	AA622344	Sequence 6513	AA716751
Sequence 6462	AA625222	Sequence 6514	AA723130
Sequence 6463	AA625682	Sequence 6515	AA723679
Sequence 6464	AA626439	Sequence 6516	AA724619
Sequence 6465	AA628526	Sequence 6517	AA724868
Sequence 6466	AA629534	Sequence 6518	AA729119
Sequence 6467	AA629608	Sequence 6519	AA732326
Sequence 6468	AA631868	Sequence 6520	AA740261
Sequence 6469	AA631903	Sequence 6521	AA741056
Sequence 6470	AA633794	Sequence 6522	AA742231
Sequence 6471	AA634039	Sequence 6523	AA744367
Sequence 6472	AA634216	Sequence 6524	AA747819
Sequence 6473	AA634379	Sequence 6525	AA764981
Sequence 6474	AA638989	Sequence 6526	AA767323
Sequence 6475	AA639317	Sequence 6527	AA772603
Sequence 6476	AA640105	Sequence 6528	AA773009
Sequence 6477	AA640141	Sequence 6529	AA773362
Sequence 6478	AA641623	Sequence 6530	AA773374
Sequence 6479	AA642471	Sequence 6531	AA775914
Sequence 6480	AA648508	Sequence 6532	AA778038
Sequence 6481	AA649193	Sequence 6533	AA778109
Sequence 6482	AA652846	Sequence 6534	AA778596
Sequence 6483	AA653353	Sequence 6535	AA808003
Sequence 6484	AA654557	Sequence 6536	AA810159
Sequence 6485	AA654675	Sequence 6537	AA812193
Sequence 6486	AA654875	Sequence 6538	AA813266
Sequence 6487	AA659719	Sequence 6539	AA813941
Sequence 6488	AA662658	Sequence 6540	AA814496
Sequence 6489	AA662702	Sequence 6541	AA826649
Sequence 6490	AA662822	Sequence 6542	AA827331
Sequence 6491	AA664480	Sequence 6543	AA831795
Sequence 6492	AA668224	Sequence 6544	AA831876
Sequence 6493	AA668692	Sequence 6545	AA835377
Sequence 6494	AA668951	Sequence 6546	AA836233
Sequence 6495	AA670280	Sequence 6547	AA836488
Sequence 6496	AA675891	Sequence 6548	AA844979
Sequence 6497	AA677282	Sequence 6549	AA845409
Sequence 6498	AA680052	Sequence 6550	AA845442
Sequence 6499	AA682910	Sequence 6551	AA846152

Sequence 6552	AA846576	Sequence 6604	AI028507
Sequence 6553	AA854071	Sequence 6605	Al028516
Sequence 6554	AA855093	Sequence 6606	Al033912
Sequence 6555	AA862250	Sequence 6607	AI038038
Sequence 6556	AA876668	Sequence 6608	A1038986
Sequence 6557	AA884922	Sequence 6609	AI039271
Sequence 6558	AA884972	Sequence 6610	A1039636
Sequence 6559	AA884992	Sequence 6611	AI040598
Sequence 6560	AA886748	Sequence 6612	AI041497
Sequence 6561	AA886957	Sequence 6613	AI042152
Sequence 6562	AA886998	Sequence 6614	AI050872
Sequence 6563	AA887054	Sequence 6615	AI051210
Sequence 6564	AA889969	Sequence 6616	AI052124
Sequence 6565	AA890326	Sequence 6617	AI052126
Sequence 6566	AA890421	Sequence 6618	AI052724
Sequence 6567	AA896959	Sequence 6619	AI052738
Sequence 6568	AA902127	Sequence 6620	A1053597
Sequence 6569	AA907003	Sequence 6621	AI054023
Sequence 6570	AA909768	Sequence 6622	AI057281
Sequence 6571	AA916753	Sequence 6623	A1057555
Sequence 6572	AA921906	Sequence 6624	AI064691
Sequence 6573	AA923172	Sequence 6625	AI074143
Sequence 6574	AA926926	Sequence 6626	AI074389
Sequence 6575	AA927818	Sequence 6627	Al075893
Sequence 6576	AA934755	Sequence 6628	AI077500
Sequence 6577	AA934851	Sequence 6629	AI079882
Sequence 6578	AA935181	Sequence 6630	Al082272
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Sequence 6580	AA935560	Sequence 6632	AI082674
Sequence 6581	AA935845	Sequence 6633	AI084100
Sequence 6582	AA937773	Sequence 6634	Al084101
Sequence 6583	AA938717	. Sequence 6635	Al088178
Sequence 6584	AA953892	Sequence 6636	AI090786
Sequence 6585	AA961085	Sequence 6637	AI093233
Sequence 6586	AA968756	Sequence 6638	AI094394
Sequence 6587	AA969214	Sequence 6639	AI095250
Sequence 6588	AA972883	Sequence 6640	Al096410
Sequence 6589	AA974621	Sequence 6641	Al123199
•	AA974908	Sequence 6642	Al123672
Sequence 6590	AA975275	Sequence 6643	Al123822
Sequence 6591		•	Al125642
Sequence 6592	AA977871	Sequence 6644	Al126257
Sequence 6593	AA984586	Sequence 6645	
Sequence 6594	AA988125	Sequence 6646	Al127268 Al127556
Sequence 6595	AA989049	Sequence 6647	
Sequence 6596		Sequence 6648	Al129800
Sequence 6597		Sequence 6649	Al131216
Sequence 6598		Sequence 6650	AI133208
Sequence 6599		Sequence 6651	Al133690
Sequence 6600		Sequence 6652	AI139036
Sequence 6601	AI016162	Sequence 6653	Al142021
Sequence 6602		Sequence 6654	Al142095
Sequence 6603	AI022058	Sequence 6655	Al142558

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Sequence 6656	Al146476	Sequence 6708	Al267849
Sequence 6657	Al148115	Sequence 6709	Al269060
Sequence 6658	Al148251	Sequence 6710	Al272941
Sequence 6659	Al148561	Sequence 6711	Al273945
Sequence 6660	Al148676	Sequence 6712	A1275792
Sequence 6661	Al167651	Sequence 6713	AI282318
Sequence 6662	AI174853	Sequence 6714	Al283096
Sequence 6663	AI174899	Sequence 6715	Al285408
Sequence 6664	Al184748	Sequence 6716	AI287603
Sequence 6665	Al189386	Sequence 6717	Al289111
Sequence 6666	Al189897	Sequence 6718	Al290826
Sequence 6667	AI190785	Sequence 6719	Al291296
Sequence 6668	AI198311	Sequence 6720	Al298972
Sequence 6669	AI198853	Sequence 6721	Al300489
Sequence 6670	Al198930	Sequence 6722	Al302102
Sequence 6671	Al198986	Sequence 6723	Al305838
Sequence 6672	Al199094	Sequence 6724	Al306457
Sequence 6673	Al199681	Sequence 6725	Al307419
Sequence 6674	Al200798	Sequence 6726	AI310138
Sequence 6675	Al200830	Sequence 6727	Al312552
Sequence 6676	Al201564	Sequence 6728	Al313387
Sequence 6677	Al203377	Sequence 6729	Al332712
Sequence 6678	Al204096	Sequence 6730	Al335829
Sequence 6679	Al206483	Sequence 6731	Al338379
Sequence 6680	Al206818	Sequence 6732	Al340611
Sequence 6681	Al207546	Sequence 6733	Al341551
Sequence 6682	Al207618	Sequence 6734	Al342409
Sequence 6683	Al207650	Sequence 6735	Al342468
Sequence 6684	Al214272	Sequence 6736	Al342669
Sequence 6685	Al216968	Sequence 6737	Al343112
Sequence 6686	Al216984	Sequence 6738	Al343692
Sequence 6687	Al217003	Saguanca 6720	Al346657
Sequence 6688	Al217172	Sequence 6740	Al348579
Sequence 6689	Al219570	Sequence 6741	Al352141
Sequence 6690	Al221894	Sequence 6742	Al357472
Sequence 6691	Al222329	Sequence 6743	Al358685
Sequence 6692	Al222789	Sequence 6744	Al360710
•	Al222709 Al222806	Sequence 6745	Al361905
Sequence 6693	Al223061	Sequence 6746	Al361969
Sequence 6694		Sequence 6747	Al365612
Sequence 6695	Al244217	•	Al376840
Sequence 6696	AI253288	Sequence 6748	Al380706
Sequence 6697	AI253319	Sequence 6749 Sequence 6750	Al380932
Sequence 6698	AI253338	•	Al382020
Sequence 6699	AI253348	Sequence 6751	
Sequence 6700	AI261883	Sequence 6752	Al382065
Sequence 6701	AI267162	Sequence 6753	Al382189
Sequence 6702	AI267353	Sequence 6754	Al383945
Sequence 6703	AI267454	Sequence 6755	Al400752
Sequence 6704	Al267519	Sequence 6756	AI401771
Sequence 6705	AI267652	Sequence 6757	Al420898
Sequence 6706	Al267656	Sequence 6758	AI423180
Sequence 6707	Al267664	Sequence 6759	Al432306

		0 0040	A1004 40E
Sequence 6760	Al432462		AI681405
Sequence 6761	Al433976		AI683298
Sequence 6762	Al435153	***************************************	Al684170
Sequence 6763	Al435945		Al686139
Sequence 6764	A1440088	Coque in the contract of the c	A1689722
Sequence 6765	Al452485	2242	A1693697
Sequence 6766	Al457157		A1694087
Sequence 6767	Al467861		Al694199
Sequence 6768	Al468347	•••	Al694767
Sequence 6769	Al474235		AI697470
Sequence 6770	Al475476	Sequence 6822	Al701140
Sequence 6771	AI523940	Sequence 6823	AI703256
Sequence 6772	AI525552	Sequence 6824	AI708172
Sequence 6773	AI525702	Sequence 6825	AI709186
Sequence 6774	AI525843	Sequence 6826	AI718780
Sequence 6775	AI536991	Sequence 6827	AI718825
Sequence 6776	AI553845	Sequence 6828	Al733116
Sequence 6777	AI557112	Sequence 6829	AI734854
Sequence 6778	AI557182	Sequence 6830	AI735314
Sequence 6779	AI557225	Sequence 6831	AI735683
Sequence 6780	AI557237	Sequence 6832	Al739102
Sequence 6781	AI557452	Sequence 6833	AI741354
Sequence 6782	AI557599	Sequence 6834	AI741843
Sequence 6783	AI557639	Sequence 6835	AI742468
Sequence 6784	AI567102	Sequence 6836	AI742912
Sequence 6785	AI569556	Sequence 6837	A1743852
Sequence 6786	AI569819	Sequence 6838	AI744509
Sequence 6787	AI569955	Sequence 6839	A1749547
Sequence 6788	AI570221	Sequence 6840	A1751364
Sequence 6789	AI571418	Sequence 6841	Al751565
Sequence 6790	AI580176	Sequence 6842	A1753664
Sequence 6791	AI590195	Sequence 6843	AI753784
Sequence 6792	AI590351	Sequence 6844	AI754741
Sequence 6793	AI598180	Sequence 6845	A1755085
Sequence 6794	Al610920	Sequence 6846	AI760827
Sequence 6795	Al620111	Sequence 6847	AI761110
Sequence 6796	Al620796	Sequence 6848	AI766123
Sequence 6797	Al623924	Sequence 6849	AI766514
Sequence 6798	Al625201	Sequence 6850	AI768465
Sequence 6799	AI625864	Sequence 6851	AI769392
Sequence 6800	Al631703	Sequence 6852	AI770054
Sequence 6801	Al631745	Sequence 6853	AI791548
Sequence 6802	Al634228	Sequence 6854	AI793062
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Sequence 6804	Al635681	Sequence 6856	A1799974
Sequence 6805	Al636899	Sequence 6857	AI800048
Sequence 6806	Al648684	Sequence 6858	AI802154
•		Sequence 6859	AI805364
Sequence 6807		Sequence 6860	AI813758
Sequence 6808		Sequence 6861	AI815380
Sequence 6809		Sequence 6862	AI819014
Sequence 6810		Sequence 6863	Al827550
Sequence 6811	Al672868	Sequence 6003	A1021330

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Sequence 6864	AI828670		AL121547
Sequence 6865	Al831047		AW003247
Sequence 6866	Al861878	Sequence 6918	AW003692
Sequence 6867	Al879179	Sequence 6919	AW005103
Sequence 6868	A1879995	Sequence 6920	AW008213
Sequence 6869	AI887632	Sequence 6921	AW015683
Sequence 6870	Al888438	Sequence 6922	AW020479
Sequence 6871	A1890240	Sequence 6923	AW022300
Sequence 6872	Al910552	Sequence 6924	AW022756
Sequence 6873	Al912414	Sequence 6925	AW023489
Sequence 6874	AI914128	Sequence 6926	AW051386
Sequence 6875	Al914133	Sequence 6927	AW052140
Sequence 6876	Al921434	Sequence 6928	AW062933
Sequence 6877	AI922855	Sequence 6929	AW067913
Sequence 6878	Al923011	Sequence 6930	AW071063
Sequence 6879	AI923978	Sequence 6931	AW071903
Sequence 6880	Al929509	Sequence 6932	AW080160
Sequence 6881	AI937296	Sequence 6933	AW083804
Sequence 6882	Al940170	Sequence 6934	AW090544
Sequence 6883	Al948503	Sequence 6935	AW104561
Sequence 6884	AI954458	Sequence 6936	AW135430
Sequence 6885	Al961840	Sequence 6937	AW138413
Sequence 6886	AI963623	Sequence 6938	AW151295
Sequence 6887	Al970594	Sequence 6939	AW157047
Sequence 6888	Al972279	Sequence 6940	AW157303
Sequence 6889	Al973109	Sequence 6941	AW160731
Sequence 6890	Al983220	Sequence 6942	AW161491
Sequence 6891	AL035731	Sequence 6943	AW168960
Sequence 6892	AL035985	Sequence 6944	AW169876
Sequence 6893	AL036123	Sequence 6945	AW170485
Sequence 6894	AL036169	Sequence 6946	AW179006
Sequence 6895	AL036483	Sequence 6947	AW183759
Sequence 6896	AL036801	Sequence 6948	C06419
Sequence 6897	AL036988	Sequence 6949	D55192
Sequence 6898	AL037267	Sequence 6950	D58694
Sequence 6899	AL037471	Sequence 6951	F05508
Sequence 6900	AL037800	Sequence 6952	F06621
Sequence 6901	AL039253	Sequence 6953	F06739
Sequence 6902	AL040873	Sequence 6954	F12493
Sequence 6903	AL041050	Sequence 6955	F22763
Sequence 6904	AL041780	Sequence 6956	H00112
Sequence 6905	AL042316	Sequence 6957	H03785
Sequence 6906	AL044447	Sequence 6958	H12527
Sequence 6907	AL044649	Sequence 6959	H13339
Sequence 6908	AL045213	Sequence 6960	H16455
Sequence 6909	AL048699	Sequence 6961	H20086
Sequence 6910	AL049012	Sequence 6962	H20888
Sequence 6911	AL110412	Sequence 6963	H22853
Sequence 6912	AL118621	Sequence 6964	H23837
Sequence 6913		Sequence 6965	H39906
Sequence 6914		Sequence 6966	H39960
Sequence 6915		Sequence 6967	H45550
Dequence 0313	/ \L \Z0070	00400.0000	

Sequence 6968	H50566	Sequence 7020	R89611
Sequence 6969	H57382	Sequence 7021	R93759
Sequence 6970	H59731	Sequence 7022	R98423
Sequence 6971	H59791	Sequence 7023	T05006
Sequence 6972	H70670	Sequence 7024	T05635
Sequence 6973	H75695	Sequence 7025	T10307
Sequence 6974	H78954	Sequence 7026	T27258
Sequence 6975	H81659	Sequence 7027	T28083
Sequence 6976	H83574	Sequence 7028	T31772
Sequence 6977	H84729	Sequence 7029	T61421
Sequence 6978	H86010	Sequence 7030	T65998
Sequence 6979	L49115	Sequence 7031	T72376
Sequence 6980	M85423	Sequence 7032	T78948
Sequence 6981	N25145	Sequence 7033	T83709
Sequence 6982	N26359	Sequence 7034	T84943
Sequence 6983	N26421	Sequence 7035	U69559
Sequence 6984	N28384	Sequence 7036	W00556
Sequence 6985	N30486	Sequence 7037	W03170
Sequence 6986	N33567	Sequence 7038	W03471
Sequence 6987	N40852	Sequence 7039	W05472
Sequence 6988	N41386	Sequence 7040	W07109
Sequence 6989	N41827	Sequence 7041	W19899
Sequence 6990	N42196	Sequence 7042	W25222
Sequence 6991	N44001	Sequence 7043	W25970
Sequence 6992	N45940	Sequence 7044	W27440
Sequence 6993	N52271	Sequence 7045	W28596
Sequence 6994	N52554	Sequence 7046	W28729
Sequence 6995	N63991	Sequence 7047	W28994
Sequence 6996	N72922	Sequence 7048	W39249
Sequence 6997	N79747	Sequence 7049	W40294
Sequence 6998	N90917	Sequence 7050	W45179
Sequence 6999	N98328	Sequence 7051	W45265
Sequence 7000	N98569	Sequence 7052	W51959
Sequence 7001	N98593	Sequence 7053	W56523
Sequence 7002	R11581	Sequence 7054	W56718
Sequence 7003	R12113	Sequence 7055	W63793
Sequence 7004	R17092	Sequence 7056	W65292
Sequence 7005	R20331	Sequence 7057	W69806
Sequence 7006	R22136	Sequence 7058	W76134
Sequence 7007	R32613	Sequence 7059	W78775
Sequence 7008	R35797	Sequence 7060	W87891
Sequence 7009	R49734	Sequence 7061	W93655
Sequence 7010	R50467	Sequence 7062	W99251
Sequence 7011	R59147	Sequence 7063	Z19471
Sequence 7012	R61601	Sequence 7064	Z30311
Sequence 7013	R69943	Sequence 7065	A06846
Sequence 7014	R70570	Sequence 7066	AB001106
Sequence 7015	R70639	Sequence 7067	AB002357
Sequence 7016	R70995	Sequence 7068	AB002365
Sequence 7017	R71431	Sequence 7069	AB002386
Sequence 7018	R74597	Sequence 7070	AB002387
Sequence 7019	R80852	Sequence 7071	AB002451

Table 4-1

Sequence 7072	AB006077	Sequence 7124	AF000982
Sequence 7073	AB007892	Sequence 7125	AF000984
Sequence 7074	AB007896	Sequence 7126	AF007142
Sequence 7075	AB007931	Sequence 7127	AF007791
Sequence 7076	AB007956	Sequence 7128	AF010233
Sequence 7077	AB011004	Sequence 7129	AF013717
Sequence 7078	AB011079	Sequence 7130	AF013759
Sequence 7079	AB011103	Sequence 7131	AF013988
Sequence 7080	AB011120	Sequence 7132	AF016270
Sequence 7081	AB014486	Sequence 7133	AF017782
Sequence 7082	AB014519	Sequence 7134	AF020038
Sequence 7083	AB014543	Sequence 7135	AF020202
Sequence 7084	AB014548	Sequence 7136	AF020351
Sequence 7085	AB014601	Sequence 7137	AF020797
Sequence 7086	AB015639	Sequence 7138	AF021819
Sequence 7087	AB016068	Sequence 7139	AF022215
Sequence 7088	AB017004	Sequence 7140	AF026939
Sequence 7089	AB018331	Sequence 7141	AF026977
Sequence 7090	AB018346	Sequence 7142	AF027824
Sequence 7091	AB019524	Sequence 7143	AF029689
Sequence 7092	AB019563	Sequence 7144	AF034091
Sequence 7093	AB019566	Sequence 7145	AF035286
Sequence 7094	AB019568	Sequence 7146	AF035293
Sequence 7095	AB020637	Sequence 7147	AF035309
Sequence 7096	AB020661	Sequence 7148	AF038451
Sequence 7097	AB020686	Sequence 7149	AF038452
Sequence 7098	AB020689	Sequence 7150	AF038957
Sequence 7099	AB020692	Sequence 7151	AF039692
Sequence 7100	AB020705	Sequence 7152	AF039918
Sequence 7101	AB021288	Sequence 7153	AF042386
Sequence 7102	AB021663	Sequence 7154	AF044321
Sequence 7103	AB023145	Sequence 7155	AF044670
Sequence 7104	AB023163	Sequence 7156	AF044671
Sequence 7105	AB023173	Sequence 7157	AF044956
Sequence 7106	AB023205	Sequence 7158	AF045167
Sequence 7107	AB023210	Sequence 7159	AF045229
Sequence 7108	AB023230	Sequence 7160	AF047020
Sequence 7109	AB026190	Sequence 7161	AF049910
Sequence 7110	AB027013	Sequence 7162	AF051323
Sequence 7111	AB028624	Sequence 7163	AF052124
Sequence 7112	AB028981	Sequence 7164	AF052129
Sequence 7113	AB029000	Sequence 7165	AF054186
Sequence 7114	AB029004	Sequence 7166	AF054187
Sequence 7115	AB032983	Sequence 7167	AF054838
Sequence 7116	AB032991	Sequence 7168	AF054990
Sequence 7117	AB033007	Sequence 7169	AF056322
Sequence 7118	AB033071	Sequence 7170	AF059524
Sequence 7119	AB033079	Sequence 7171	AF060219
Sequence 7120	AB033091	Sequence 7172	AF061258
Sequence 7121	AB033899	Sequence 7173	AF062323
Sequence 7122	AD001528	Sequence 7174	AF065388
Sequence 7123	AF000381	Sequence 7175	AF067171

Sequence 7176	AF068235	Sequence 7228	AF131758
Sequence 7177	AF070523	Sequence 7229	AF131763
Sequence 7178	AF070539	Sequence 7230	AF131814
Sequence 7179	AF070555	Sequence 7231	AF131838
Sequence 7180	AF070556	Sequence 7232	AF131856
Sequence 7181	AF070561	Sequence 7233	AF132000
Sequence 7182	AF070650	Sequence 7234	AF141968
Sequence 7183	AF070655	Sequence 7235	AF147330
Sequence 7184	AF071202	Sequence 7236	AF147331
Sequence 7185	AF073475	Sequence 7237	AF147339
Sequence 7186	AF075587	Sequence 7238	AF147395
Sequence 7187	AF077030	Sequence 7239	AF151103
Sequence 7188	AF077035	Sequence 7240	AF151109
Sequence 7189	AF077042	Sequence 7241	AF151803
Sequence 7190	AF077202	Sequence 7242	AF151818
Sequence 7191	AF078848	Sequence 7243	AF151840
Sequence 7192	AF078849	Sequence 7244	AF151867
Sequence 7193	AF078854	Sequence 7245	AF151871
Sequence 7194	AF078859	Sequence 7246	AF151878
Sequence 7195	AF078862	Sequence 7247	AF153608
Sequence 7196	AF082657	Sequence 7248	AF155095
Sequence 7197	AF084457	Seguence 7249	AF155110
Sequence 7198	AF084555	Sequence 7250	AF156965
Sequence 7199	AF085355	Sequence 7251	AF159056
Sequence 7200	AF085845	Sequence 7252	AF168956
Sequence 7201	AF086002	Sequence 7253	AF169797
Sequence 7202	AF086003	Sequence 7254	AF170583
Sequence 7203	AF086172	Sequence 7255	AF174605
Sequence 7204	AF086182	Sequence 7256	AF176574
Sequence 7205	AF086183	Sequence 7257	AF179274
Sequence 7206	AF086336	Sequence 7258	AF182289
Sequence 7207	AF086401	Sequence 7259	AF182645
Sequence 7208	AF086431	Sequence 7260	AF188745
Sequence 7209	AF086517	Sequence 7261	AF188746
Sequence 7210	AF086557	Sequence 7262	AF195120
Sequence 7211	AF087999	Sequence 7263	AF201077
Sequence 7212	AF089747	Sequence 7264	AJ002744
Sequence 7213	AF092092	Sequence 7265	AJ004832
Sequence 7214	AF097514	Sequence 7266	AJ005016
Sequence 7215	AF100741	Sequence 7267	AJ010071
Sequence 7216	AF100749	Sequence 7268	AJ011001
Sequence 7217	AF100756	Sequence 7269	AJ223183
Sequence 7218	AF100757	Sequence 7270	AJ249248
Sequence 7219	AF102265	Sequence 7271	AL021683
Sequence 7220	AF103907	Sequence 7272	AL035304
Sequence 7221	AF109219	Sequence 7273	AL049381
Sequence 7222	AF113131	Sequence 7274	AL049705
Sequence 7223	AF113132	Sequence 7275	AL049932
Sequence 7224	AF124438	Sequence 7276	AL049934
Sequence 7225	AF126780	Sequence 7277	AL049957
Sequence 7226	AF126782	Sequence 7278	AL049969
Sequence 7227	AF129927	Sequence 7279	AL050159
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Table 4-1

Sequence 7280	AL050162	Sequence 7332	D50931
Sequence 7281	AL050187	Sequence 7333	D59253
Sequence 7282	AL050198	Sequence 7334	D78275
Sequence 7283	AL050265	Sequence 7335	D79205
Sequence 7284	AL050287	Sequence 7336	D79986
Sequence 7285	AL079298	Sequence 7337	D80009
Sequence 7286	AL080202	Sequence 7338	D83780
Sequence 7287	AL080209	Sequence 7339	D84145
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Sequence 7295	AL117423	Sequence 7347	D87742
Sequence 7296	AL117429	Sequence 7348	D88208
Sequence 7297	AL117446	Sequence 7349	D89053
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Sequence 7299	AL117621	Sequence 7351	D90373
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Sequence 7301	AL117666	Sequence 7353	E02628
Sequence 7302	AL122088	Sequence 7354	E02822
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Sequence 7305	D00017	Sequence 7357	E08663
Sequence 7306	D12676	Sequence 7358	J02876
Sequence 7307	D13119	Sequence 7359	J02888
Sequence 7308	D13315	Sequence 7360	J03007
Sequence 7309	D13866	Sequence 7361	J03040
Sequence 7310	D14530	Sequence 7362	J03248
Sequence 7311	D14696	Sequence 7363	J03464
Sequence 7312	D14697	Sequence 7364	J03746
Sequence 7313	D14812	Sequence 7365	J03799
Sequence 7314	D16911	Sequence 7366	J04164
Sequence 7315	D17039	Sequence 7367	J04205
Sequence 7316	D17554	Sequence 7368	J04208
Sequence 7317	D21262	Sequence 7369	J04443
Sequence 7318	D21853	Sequence 7370	J04794
Sequence 7319	D25542	Sequence 7371	K01911
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Sequence 7321	D28476	Sequence 7373	L00635
Sequence 7322	D29677	Sequence 7374	L01100
Sequence 7323	D29954	Sequence 7375	L04636
Sequence 7324	D30655	Sequence 7376	L05093
Sequence 7325	D31883	Sequence 7377	L05779
Sequence 7326	D37931	Sequence 7378	L08441
Sequence 7327	D38293	Sequence 7379	L09159
Sequence 7328	D38441	Sequence 7380	L10320
Sequence 7329	D38583	Sequence 7381	L10678
Sequence 7330	D42047	Sequence 7382	L16558
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Table 4-1

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Sequence 7386	L20941	Sequence 7438	M76729
Sequence 7387	L26081	Sequence 7439	M77830
Sequence 7388	L33404	Sequence 7440	M80783
Sequence 7389	L34840	Sequence 7441	M81757
Sequence 7390	L38486	Sequence 7442	M83822
Sequence 7391	L38961	Sequence 7443	M90054
Sequence 7392	L47276	Sequence 7444	M93651
Sequence 7393	L49504	Sequence 7445	M95724
Sequence 7394	L77964	Sequence 7446	M97347
Sequence 7395	M10119	Sequence 7447	S66196
Sequence 7396	M10941	Sequence 7448	S66431
Sequence 7397	M11058	Sequence 7449	S67815
Sequence 7398	M11353	Sequence 7450	S69272
Sequence 7399	M12938	Sequence 7451	S70154
Sequence 7400	M14200	Sequence 7452	S70290
Sequence 7401	M14631	Sequence 7453	S73498
Sequence 7402	M15178	Sequence 7454	S73591
Sequence 7403	M16768	Sequence 7455	S78203
Sequence 7404	M16804	Sequence 7456	S82081
Sequence 7405	M16942	Sequence 7457	U03886
Sequence 7406	M17733	Sequence 7458	U05875
Sequence 7407	M17885	Sequence 7459	U08815
Sequence 7408	M21154	Sequence 7460	U09559
Sequence 7409	M21574	Sequence 7461	U09820
Sequence 7410	M21895	Sequence 7462	U13991
Sequence 7411	M24194	Sequence 7463	U14658
Sequence 7412	M24902	Sequence 7464	U14967
Sequence 7413	M25077	Sequence 7465	U14970
Sequence 7414	M25246	Sequence 7466	U14971
Sequence 7415	M26663	Sequence 7467	U15008
Sequence 7416	M27332	Sequence 7468	U18291
Sequence 7417	M27335	Sequence 7469	U18297
Sequence 7418	M28211	Sequence 7470	U20157
Sequence 7419	M29366	Sequence 7471	U21858
Sequence 7420	M31630	Sequence 7472	U24105
Sequence 7421	M32790	Sequence 7473	U25789
Sequence 7422	M33308	Sequence 7474	U30826
Sequence 7423	M34840	Sequence 7475	U31905
Sequence 7424	M35252	Sequence 7476	U32944
Sequence 7425	M36341	Sequence 7477	U33821
Sequence 7426	M36634	Sequence 7478	U34252
Sequence 7427	M37716	Sequence 7479	U36764
Sequence 7428	M55265	Sequence 7480	U37143
Sequence 7429	M55409	Sequence 7481	U37230
Sequence 7430	M58525	Sequence 7482	U37689
Sequence 7431	M61831	Sequence 7483	U38894
Sequence 7432	M63573	Sequence 7484	U39400
Sequence 7433	M65217	Sequence 7485	U40272
Sequence 7434	M69238	Sequence 7486	U40763
Sequence 7435	M74091	Sequence 7487	U47742

Sequence 7488	U54558	Sequence 7540	X73902
Sequence 7489	U58855	Sequence 7541	X74801
Sequence 7490	U59435	Sequence 7542	X75593
Sequence 7491	U61083	Sequence 7543	X75861
Sequence 7492	U65090	Sequence 7544	X76538
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Sequence 7494	U70730	Sequence 7546	X80754
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Sequence 7501	U89436	Sequence 7553	X84908
Sequence 7502	U90904	Sequence 7554	X85129
Sequence 7503	U96759	Sequence 7555	X85372
Sequence 7504	U96915	Sequence 7556	X87176
Sequence 7505	U97519	Sequence 7557	X93036
Sequence 7506	V00594	Sequence 7558	X94323
Sequence 7507	X01630	Sequence 7559	X97065
Sequence 7508	X01742	Sequence 7560	X98743
Sequence 7509	X02530	Sequence 7561	Y00281
Sequence 7510	X02761	Sequence 7562	Y00282
Sequence 7511	X04098	Sequence 7563	Y00503
Sequence 7512	X04470	Sequence 7564	Y08319
Sequence 7513	X05332	Sequence 7565	Y10313
Sequence 7514	X06617	Sequence 7566	Y13286
Sequence 7515	X07819	Sequence 7567	Z11241
Sequence 7516	X12597	Sequence 7568	Z14244
Sequence 7517	X12791	Sequence 7569	Z18538
Sequence 7518	X15187	Sequence 7570	Z26317
Sequence 7519	X15729	Sequence 7571	Z29330
Sequence 7520	X15759	Sequence 7572	Z47087
Sequence 7521	X52174	Sequence 7573	Z74615
Sequence 7522	X53605	Sequence 7574	N91258
Sequence 7523	X55654	Sequence 7575	N91825
Sequence 7524	X56999	Sequence 7576	N92483
Sequence 7525	X57352	Sequence 7577	Q13363
Sequence 7526	X57766	Sequence 7578	Q46089
Sequence 7527	X57812	Sequence 7579	Q65520
Sequence 7528	X59406	Sequence 7580	Q65676
Sequence 7529	X59834	Sequence 7581	Q75166
Sequence 7530	X59841	Sequence 7582	Q94780
Sequence 7531	X63432	Sequence 7583	T59274
Sequence 7532	X64330	Sequence 7584	V03517
Sequence 7533	X65644	Sequence 7585	V29653
Sequence 7534	X66276	Sequence 7586	V32447
Sequence 7535	X67698	Sequence 7587	V33197
Sequence 7536	X69117	Sequence 7588	V34263
Sequence 7537	X70326	Sequence 7589	V34271
Sequence 7538	X73114	Sequence 7590	V43612
Sequence 7539	X73608	Sequence 7591	V58360

Sequence 7592	V58628	Sequence 7644	AF103908
Sequence 7593	V58648	Sequence 7645	AL022097
Sequence 7594	V58684	Sequence 7646	AL031346
Sequence 7595	V62427	Sequence 7647	AL034555
Sequence 7596	V62428	Sequence 7648	AL079338
Sequence 7597	V62430	Sequence 7649	AP000527
Sequence 7598	V70354	Sequence 7650	S65371
Sequence 7599	V70355	Sequence 7651	U15177
Sequence 7600	V87930	Sequence 7652	Z83843
Sequence 7601	X03841	Sequence 7653	Z93020
Sequence 7602	X04316		
Sequence 7603	X24060		
Sequence 7604	X26850		
Sequence 7605	X27262		
Sequence 7606	X37486		
Sequence 7607	X37532		
Sequence 7608	X39868		
Sequence 7609	X40372		
Sequence 7610	X51668		
Sequence 7611	X52235		
Sequence 7612	X59352		
Sequence 7613	X91484		
Sequence 7614	Z13088		
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Sequence 7616	Z15149		
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Sequence 7632	AC34190		
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Sequence 7634	AC004130		
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Sequence 7636	AC004456		
Sequence 7637	AC004499		
Sequence 7638	AC005004		
Sequence 7639	AC005037		
Sequence 7640	AC006057		
Sequence 7641	AC006257		
Sequence 7642	AC006285		
Sequence 7643	AF064859		

Sequence 1953: found in patent publication W099/64594 CCCTTAGCGTGGTCGCGGCCGAGGTACTCTTTCTCTCCCCTCCTCTGAATTTAATTCTTT CAACTTGCAATTTGCAAGGATTACACATTTCACTGTGTATATTGTGTTGCAAAAAAA AAAAAGTGTCTTTGTTTAAAATTACTTGGTTTGTGAATCCATCTTGCTTTTTCCCCATTG GAACTAGTCATTAACCCATCTCTGAACTGGTAGAAAAACATCTGAAGAGCTAGTCTATCA GCATCTGACAGATGAAATTGGATGGTTCTCAGAACCATTTCACCCAGACAGCCTGTTTCT ATCCTGTTTAATAAATTAATTTTTTGGGGTTCCTCTACCATGCCATAGCAAAACCCCCTGC TTCCCAAATTCTTGTCAACAATTAAAAAAGTCTGNTGGACCTTTGGAAGTTT

Sequence 1954: found in patent publication W099/40189 TTTTCCNTTTTTTTTTTTTTTTTTTTTTTTTTTTCCCCNCCAAGCTTTATCTAGNCTTAGACT TTTTAAAAAAGTTTGGGGGCAGATTCTGAATTGGCTAAAAGACATGCATTTTTAAAACTA GCAACTCTTATTTCTTTCCTTTAAAAAATACATAGCATTAAATCCCAAATCCTATTTAAAG

Sequence 1956:found in patent publication W099/64594 CTCGAGCCGGCTCGCCAGTGTGATGGGATATCTGCAGAATTCGCCCTTAGCCGTGGTCGC GGCCCGATGTA

Sequence 1958: found in patent publication W099/64594 CCCTTTCGAGCGCCCCGGGCAGGTACTATATGGACACCAGCATGGAGCAGAAAATGA GGGCAGCAGTGTCCAATCTCATTCCAGGTGAGAAGTTGCACAGTGTCCAATAGGTGCATA CATCTCCTTAGTAAGTTGTGTATTAACAATGAAATAGAAATGAAAATATATTTTTT ATTTATGTGTATTATATTTTTCAGGCAGCTAATAAGTTGGTAGGACATAATATTTAATTC GTTGGGGACCTAATTATTTATAAATTGAATGGTTAGATATTTCTTTTGGCCTAAGCCACC ATGAAAAAATCACTGAGGCGCTAAGGGAAACCATGAACTAAGAAGCCCTTCTGAGTCTCTG TTTTCTCAACTGTAAGATGAAGAAACCTGCTCCACCTTCCTCTTGCGATTGTCGAACGGC AGTTGAGATACTGCGCCAGAATGGACCCTTATTGATGGCCTAC

Sequence 1960: found in patent publication W099/64594 NGATATCTGCANAATTCGCCCTTTCGAGCGGCCGCCCGGGCAGGTACAACTTTTTAAATA GGGAATATGATAGCTTNGCATGGTGGTGTGCACCTATAGCCCCCACTGCCTGGAAAGCTG

GGGTGGGAGAATCGCTTGAGTCCAGGAGTTTGAGGTTACAGTGATCCACGATCNNGCCAC TACACTCCAGCCTGGGCANCAGAGCAAGACCCTGTNTCAAAGCATAAAATGGAATNACAT ATCAAATGAAACANGGAAAATGAAGCTGA

Sequence 1962:found in patent publication W099/64594 GCATGCTCGAGCCGGCCAGTGTGATGGGATATCTGCAGAATTCGCCCTTTCGAGCGG CCCGCCCGG

Sequence 3724: found in patent application W099/64594 CCCTTTCGAGCGCCCCGGGCAGGTACTCAGACAAATACAACGTGCCCATTTCGTCAG ATATTGCGCAGAACCAAGAATTTTATAAGAACGCAGAAGTTAGACCACCATTTACCGTAT GCATNTTTAATTAAGGCAGGCCATTCTCGAATCTCCAGAAAAGCAGCTAACACTAAATGA GATCTATAACTGGTTCACACGAATGTTTGCTTACTTCCGACGCAACGCGGCCACGTGGAA GAATGCAGTGCGTCATAATCTTAGTCTTCACAAGTGTTTTGTGGCGAGTAGAAAACGTTA AAGGGGGCAGTATGGGACAGTGGATTGAAGTANGAATTCCAAAAACGAAGGGCCCAAAAA GGATCAGTGGGTAACCCTTT

GTGGGTGGGCCAGCACCTAGAAGACACATGGTNGGTTGTGCAACATAAAATCCCCT TCAAGCGGACTCAGTGG

Sequence 3729: found in patent application W099/64594 CCCTTAGCGTGGCCGGCCGAGGTACAGAATTTATTATGAAATAGCTTAATGGCAAGTG GTAATTTAGAAGAATTAACATTAACATTAACATTAGAAATTTGAT ATATTCTTGAAGCCCTTTTACACAAGTAATTTCTATAATTTGATTGTAATGAAAGTATAA TATACCTTGTTACTATTATCAGGAATTAATTTTTTGAAAGTAGAATTCCTTAATCAAGCCAA GGTTAATGCTGCTTTATAGGAAATTAATCAGGTAGTTTAACACTAGAGCTCATTAGCCCA ACCTGTATGTAGCCAAAAAATAAATCATTCTCTGATAAATTACCNTATTAAAATAATTTTTTAATTCATACCTTTTTAAAATAA

GAACCGGGGTTATTGGCCATAAATGGGNCCCCCAAATGGGTAANCCAAACTTTAAAAAAN ATGANCCGAAANCAAAGGGNAAAATCTGGGTGGCCCTTCCATTTNATTNTNGTANNNCAA AGGGAACCTGGNNNAAGGGGGGNCCTTTTNAAAAAAA

Sequence 4148:found Patent application W099/64594 CCCTTAGCGTGGTCGCGGCCGAGGTACCACAAAAACAGAATAATTTTGAAGTTTTAGAAT

Sequence 7622: found in patent application W099/46374

Sequence 7623: found in patent application W099/51732
CGAGGTACGTTCTTCTCCAGGTTTCTCTACCCTTAGCTTATCCTCACCCTCATTTTGC
CCTGCTGGCTGCTCAAACTATGCAACAGATTCGGCATCCTCGCTTACCCATGGCCCAGTT
TGGAGGAACCTTCTCACCTTCTCCTAACACATGGGGACCATTCCCAGTGAGACCTGTGAA
TCCTGGCAACACAAATAGCTCTCCAAAGCATAATAACACAAGCCGTCTACCTAACCAGAA
CGGGACTGTTTTACCCTCAGAGTCTGCTGGACTAGCTACTGCCAGTTGTCCTATCACTGT
CTCTTCTGTAGTTGCTGCCAGTCAGCAACTGTGTCCACTAATACCCGGACTCCTTCATC
AGTCAGAAAGCAGTTGTTTGCCTGTGTGCCTAAGACAAGTCCTCCAGCAACAGTGATTTC
TTCTGTGACAAGCACTTGTAGTTCCCTGCCTTCTGTCTCCTCTGCACCTATCACTAGCGG
GCAAGCTCCCACCACATTTCTACCTGCAAGTACCTGCCCG

Sequence 7624: found in patent application W099/42470 CCGCGGTGGCCGCCCGGGCAGGTACCGATACAGAAGAAGCTGCGTTTTGAAGACACC CTGGAGTTTGTAGGGTTTGATGCGAAGATGGCTGAGGAATCCTCCTCCTCCTCATCTCACCAACTGCTGCAACATCTCAGCAGCAGCAACTTAAAAATAAGAGTATATTAATC TCTTCTGTGGCTTCGGTGCATCATGCAAACGGCCTAGCCAAATCTTCTACCACCGTCTCT AGCTTTGCTAACAGCAAACCTGGCTCTGCTAAGAAGTTAGTGATCAAGAACTTTAAAGAT AAGCCTAAATTACCAGAAAACTACACAGATGAAACCTGGCAAAAACTGAAAGAAGCAGTG GAAGCTATTCANAATAGTACCT

Sequence 7626: found in patent application W099/57132 CGCCCGGGGCAGGTACCTTAGTGAGGCTCAAAAGGATTCTTTTGGGTCTATTTTACGCCT TATCTTTGAAATTCACCACTCGGGTGAGAAAGGTGACATTGTAGTCTTTCTGGCCTGTGA ACAAGATATTGAGAAAGTCTGTGAAACTGTCTATCAAGGATCTAACCTAAACCCAGATCT TGGAGAACTGGTGGTTGTTCCTTTGTATCCAAAAGAAAATGTTCATTGTTCAAGCCACT CGATGAAACAGAAAAAAGATGCCAAGTTTATCAAAGAAGAGTGGTGTTAACTACTACTC TGGAGAGTTTTTTGATCTGGAGCAACTCAGTCAGATTTGTTATCGATGTGGGTGTGGAAAA GAAAAAAGGTGTCCTCGGC

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Table 5-1

Order	Clone	Gen Bank Accession Number
1	28106	R40780
2	30148	R41329
3	37408	R49695
4	37598	R51080
5	38554	R49731
6	40104	R54594
7	41850	R52786
8	42747	R61847
9	46365	H09774
10	46477	H09966
11	46694	H10192
12	47400	H10413
13	47652	H11433
14	50080	H17934
15	50486	H17618
16	50935	H18626
17	51378	H24011
18	51799	H23524
19	67237	T52700
20	69002	T54298
21	69935	T48692
22	81413	T60161
23	111812	T91225
24	120528	T95320
25	124447	R01094
26	127524	R08871
27	133864	R28660
28	136605	R35051
29	149288	R82644
30	151055	H02231
31	156962	R74321
	196435	R91517
32	197651	R94504
33	200354	R96914
34	204483	H58234
35	205497	H57857
36		H66840
37	210697	H80063
38	249070	N30680
39	258300	N32226
40	258649	
41	259950	N32587
42	266343	N26556
43	277173	N34316
44	278729	N62936
45	280375	N47113
46	293745	N65950

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17	321834	W37112
18	344126	W73753
19	344190	W69741
50	344721	W74673
51	346308	W74133
52	396350	AA758375
53	399101	AA733027
54	413148	AA707871
55	415473	W80447
56	415610	W80730
57	415619	W80741
58	417393	W88942
59	418150	W90164
60	418206	W90529
61	418279	W90323
62	428560	AA005401
63	431408	AA706839
64	432479	AA699494
65	449428	AA777883
66	451596	AA707080
67	451911	AA706969
68	455275	AA677575
69	490484	AA101632
70	501431	· AA115300
71	503741	AA131466
72	595318	AA164301
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74	645565	AA204830
75	687297	AA235116
76	712950	AA282273
77	729953	AA412049
78	731218	AA416724
79	742049	AA401457
80	742569	AA400074
81	753745	AA410491
82	755304	AA436327
83	767289	AA418418
84	767823	AA418728
85	768217	AA424905
86	782233	AA431721
87	782503	AA431773
88	785663	AA449085
89	786525	AA452113
90	786602	AA478470
91	788180	AA453310
92	795499	AA454215
92	810873	AA459197
94	810987	AA485357
94 95	815183	AA481152
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96	815556	AA456821
97	824723	AA488986
98	839081	AA487608
99	840878	AA482324
100	841141	AA487031
101	845345	AA773478
102	884606	AA630006
103	897924	AA598808
104	950470	AA599102
105	1030855	AA621761
106	1049033	AA778675
107	1049185	AA620697
108	1049230	AA620715
109	1292432	AA718910
110	1323448	AA873604
111	1412481	AA845156
112	1416782	AA894557
113	1420370	AA857035
114	1468220	AA884897
115	1471841	AA873355
116	1472753	AA872402

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Table 5-2

Order	Clone	Gen Bank Accession Number
4	629896	AA219045
1	897567	AA489611
2	856650	AA669314
3	897781	AA598517
4	882510	AA676460
5	586685	AA130579
6	769890	AA430382
7	525926	AA074511
8	714414	AA293215
9	812251	AA455056
10	192569	H41489
11	•	AA465723
12	814989	AA430032
13	781089	AA426113
14	757265 359661	AA011096
15	768562	AA429297
16	811062	AA485441
17	878676	AA775355
18		AA478298
19	740941	AA251770
20	684655	AA489261
21	842863	AA670408
22	878798	N47691
23	281010	R32952
24	135221	AA150500
25	491751	AA434403
26	770879	N71982
27	290841	N72263
28	291341	N50880
29	281003	AA476576
30	785368	H08120
31	45582	AA478066
32	739511	AA427570
33	770043	AA187351
34	624627	W86653
35	416833	AA063521
36	359982	AA253464
37	669375	AA629262
38	744047	AA706968
39	451907	AA426025
40	757244	AA700904
41	453107	AA419177
42	755578	AA156802
43	502518	AA156602 AA425299
44	773286	
45	843321	AA489569
46	278687	N62924

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7	853006	AA668230
8	897529	AA497001
9	757165	AA443950
50	230971	R96155
51	796624	AA460530
52	38740	R51273
53	868757	AA775325
54	825372	AA504507
55	746245	AA417761
56	292399	N68399
57	626716	AA191548
58	489106	AA056538
59	22074	T66180
60	85800	T72067
61	705064	AA279990
62	277740	N49589
63	35300	R43798
64	788256	AA454098
65	746007	AA482007
66	455263	AA677572
67	277112	N39611
68	259591	N32768
69	22773	R38613
70	269997	N24910
71	451918	AA706964
72	345626	W72051
73	51041	H18633
74	32299	R42685
75	812266	AA455062
76	41391	R56123
77	878253	AA775791
77 78	83920	T64312
79	740604	AA479795
80	770066	AA430545
81	196189	R92281
82	46367	H09959
83	784772	AA478542
84	246430	N53031
85	310493	W31074
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86	459941	AA779380
87	200814	R98851
88	415102	W95001
89	489677	AA099568
90	193892	H51765
91	359781	AA011320
92		AA455062
93	812266	N34436
94	277414	H15533
95	49284	1110000

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96	66317	T66816
97	239611	H79534
98	502151	AA129777
99	773330	AA425450
100	266819	N24113
101	29063	R40970
102	785293	AA476543
103	824933	AA489040
104	345616	W72431
105	144816	R76281
106	291985	N73101
107	256907	N30096
108	487793	AA043501
109	840776	AA486082
110	199945	R97066
111	232946	H75599
112	197525	H52001
113	753467	AA406551
114	771303	AA443638
115	50879	H18424
116	341680	W60414
117	461727	AA682293
118	345793	W70128
119	486591	AA042990
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121	452374	AA700876
122	241699	H91641
123	262060	H99075

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Table 6

ACC. NO.	GI NUMBER
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AA001219	1437294
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AA001734	1445600
AA001745	1445539
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Table 6

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AA007529	1463505
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AA009609	1470750
AA009611	1470820
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AA009803	1470587
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AA009840	1470887
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AA009878	1470905
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AA010083	1471131
AA010091	1471275
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AA010136	1471164
AA010167	1471343
AA010174	1471350
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AA010872	1471918
AA010932	1471959
AA010981	1472008
AA010986	1472013
AA011000	1472027
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	1472704
AA011597	1472658
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AA013018	1474045
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AA018980	1482372
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AA019316	1482727
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AA021434	1485150
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AA024967	1489872
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AA026399	1492300
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AA026609	1492465
AA026630	
AA026658	1492723
AA026682	1492849
AA026715	1492496
AA026797	1493134
AA026983	1493174
AA027037	1493227
AA027090	1493299
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AA027862	1493949
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AA028171	1494296
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AA028987	1496389
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AA029157	1496559
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AA029403	1497030
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AA029428	1496957
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AA029584	1496988
AA029673	1497076
AA029676	1497079
AA029722	1497135
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AA029881	1496108
AA029889	1496145
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AA029997	1496428
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AA030046	1496272
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AA031293	1501246
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AA031762	1501716
AA031795	1501730
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AA032221	1502183
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AA037217	1512342
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AA037450	1512752
AA037483	1512582
AA037554	1512654
AA037574	1512692
AA037620	1512756
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AA039504	1515800
AA039506	1515802
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AA039734	1516236
AA039750	1516029
AA039807	1516296
AA039812	1516301
AA039851	1516129
AA039967	1516280
AA039998	1516275
AA040037	1516380
AA040127	1516405
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AA040140	1516418
AA040180	1516596
AA040209	1516485
AA040265	1516670
AA040332	1516663
AA040405	1516766
AA040424	1516702
AA040437	1516715
AA040439	1516717
AA040499	1516786
AA040617	1517030
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AA040624	1517103
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AA040673	1516951
AA040694	1517008
AA040710	1516988
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AA040861	1517157
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AA041250	1517484
AA041467	1517765
AA041494	1517728
AA041499	1517733
AA042812	1522467
AA042813	1522468
AA042858	1522374
AA042928	1522610
AA042990	1522505
AA043109	1520983
AA043133	1521126
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AA043347	1521421
AA043415	1521290

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<u>ACC. NO.</u> AA043431	1521354
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AA044187	1522044
AA044209	1522066
AA044239	1522115
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AA301408	1953742
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AA301907	1954240
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AA304827	1957382
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AA305627	1957952
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AA314872	1967221
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AA315069	1967408
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AA317659	1969985
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AA330234	1982562
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AA330883	1983125
AA331359	1983602
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AA332593	1985086
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AA332716	1984980
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AA333358	1985601
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AA334452	1986696
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AA344393	1996703
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AA344743	1996980 1997265
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AA345762	1998186
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AA374222	2026715
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MA3 / 004 /	2023104

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AA896959	3033352
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AA902388	3037511
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AA902721	3037844
AA902831	3037954
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AA909768	3050567
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AA911154	3050444
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AA911317	3050681
AA911903	3051295
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AA916635	3056027
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AA918023	3057913
AA918370	3058260
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AA922237	3069546
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AA922326	3069635
AA923172	3070481
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AA962587	3134751
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AA972602	3147782
AA972625	3147805
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AA973273	3148453
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AA973927	3149107
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AA983362	3161887
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AA984134	3162659
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AA984744	3163269
AA985190	3163715
AA985327	3163852
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AA987644	3173008
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AA988102	3173466
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AA988468	3174160
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AA994966	3181455
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AA995066	3181555
AA995483	3181972
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AA996028	3182517
AA996104	3182593
AA996131	3182620
AA996230	3182719
AA99953	3190508
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AB000220	1843400
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AB000509	1813326
AB000584	2189952
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AB001106	1850083
AB001636	2696612
AB001872	2879897
AB002303	2224550
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AB004304	4176417
AB004788	3184185
AB004851	3702685
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AB005289	3228278
AB005535	2326258
AB005621	2832907
AB006077	2564010
AB006534	2924619
AB006621	2564313
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AB006968	2463524
AB007191	8698842
AB007510	2463576
AB007618	2465177
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AB007873	2887418
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AB007893	2887436
AB007896	2662152
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AB007916	6683704
AB007929	3413881
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AB011004	3273315
AB011079	2897815
AB011087	3043553
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AB011472	4519430
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AB014547	3327107
AB014548	3327109
AB014552	3327117
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AB015907	3721863
AB016068	5821142
AB016087	
AB016533	4586430
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AB017007	4519442
AB017430	4520349
AB017971	3978556
AB018080	3882148
AB018257	3882166
AB018266	3882170
AB018268	3882174
AB018270	3882182
AB018274	3882186
AB018276	3882194
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AB018281	3882202
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AB018296	3882236
AB018301	3882242
AB018304 AB018305	3882244
AB018313	3882260
AB018315	3882264
AB018319	3882272
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AB018324	3882282
AB018327	3882288
AB010327	3882294
AB018331	3882296
AB018334	3882302
AB018337	3882308
AB018338	3882310
AB018340	3882314
AB018342	3882318
AB018344	3882322
AB018346	3882326
AB018351	3882336
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AB018358	6009489
AB019002	4587082
AB019392	4587122
AB019435	4760646
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AB019691	5051742
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AB020631	4240136

Table 6

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AB020686	4240252
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AB020692 AB020693	4240260
AB020693	4240262
AB020698	4240270
AB020705	4240284
AB020707	4240288
AB020723	4240320
AB020724	4240322
AB020866	4003386
AB020880	4996281
AB020981	4996287
AB021179	4062855
AB021288	4038732
AB021663	4996450
AB021981	4903003
AB022663	5019617
AB023145	4589487 4589509
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AB023153	4589523
AB023157	4589535
AB023163 AB023173	4589555
AB023173 AB023177	4589563
AB023177 AB023179	4589567
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AB023102 AB023196	4589601
AB023198	4589605
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AB023205	4589619
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AB023210	4589629
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AB024334	6016837 4520137
AB024435	4520137
AB024518	4320327

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AB026723	5931601
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AB027466	6172220
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AB028945	5689380
AB028948	5689386
AB028951	5689392
AB028956	5689402
AB028964	5689418
AB028969	5689428
AB028971	5689432
AB028973	5689436
AB028977	5689444
AB028981	5689452
AB028995	5689480
AB029000	5689490
AB029004	5689498
AB029008	5689506
AB029020	5689530
AB029025	5689540
AB029027	5689544
AB029028	5689546
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AB029290	5821433
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AB032990	6330170
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AB033029	6330432
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AB033080	6331191
AB033091	6331191
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AB033114	6331406
AB033899	6174679
AC000065	1669367
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AC002040	2347081
AC002064	2076723
AC002086	2085785
AC003991	2772535
AC004022	3598727
AC004130	2842785
AC004223	3253129
AC004456	2979597
AC004499	2996641
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AC004656	3253114
AC004706	3176728
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AC005004	6624123
AC005037	4827310
AC005154	3242763
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AC005480	4835818
AC005538	4508129
AC005831	4165331
AC005880	6249667
AC005881	6382477
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Table 6

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Н54752	995172
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Н60247	1013079
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H75599	1049527
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Table 6

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Н84759	1064067
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Н85806	1067385
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н91/21 Н92106	1087684
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H92216	1087794
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Н92683	1099011
H92821	1099149
	

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Н93381	1099709
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Н95633	1108775
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H96504	1109972
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Н96643	1110129
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Н96908	1110394
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н97090	1114070
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Н97186	1114229
Н97488	1118373
Н97508	1118393
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Н97778	1118663
Н97851	1118736
Н97880	1118765
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Н98688	1123356
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J00194	188231
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J04478	179697 339666
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J04621 J04759	190280
J04794	178480 1311703
J04823 J04970	5809681
J04973	180927 3 4 0216
J05021 J05032	179101
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J05192 J05211	181607
J05480	179807 189856
J05533 J05594	1203981
J05633 K00409	186504 188523
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K03002 K03515	180468 189237
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L00635 L01042	292032 184097
L01100	184507
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L07540	190153
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L20941	507251
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U37143	1185451
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U40763	1117967
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U41813	1184168
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U42412	1335855
U42594	4096861
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U51920	1256819
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U54831	1354506
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U78575	1743870
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U79273	1710239
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U80456	2062416
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U81006	1737489
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U82226	1848270
U82256	1773058
U82756	2853276
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U83857	2623760
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U96759	2738243
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W01211	1273190
W01739	1273719
W01800	1274001
W02073	1274053
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W02333	1274331
W02342	1274349
W02426	1274612
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W02575	1274859
W02978	1274985
W03029	1275007
W03170	1275188
W03191	1275209
W03366	1275317
W03386	1275249
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W04191	1276100
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W04380	1276288
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W04588	1277455
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W15477	1289996
W15554	1289984
W16424	1289598
W16659	1291258
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W16832	1291220
W19519	1295437
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W19899	1295768
W19970	1295840
W20438	1295055
W20458	1295228
W20462	1295061
W20486	1295075
W21047	1297923
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W21894	1298776
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W27440	1307307
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W32303	1313314
W32408	1313398
W32729	1313720
W35236	1317142
W35301	1317207
W35398	1317334
W37112	1318961
W37306	1319037
W37322	1318917
W37343	1318976
W37418	1319012
W37447	1319061
W37532	1319146
W37602	1319216
W37614	1319228
W37753	1319347
W37778	1319589
W37779	1319590
W37862	1319526
W37945	1319539
W38407	1320008
W38679	1320384
W38726	1320433
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W45330	1329422
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W47364	1332003
W47487	1332274
W47576	1332227
W47585	1332236
W48700	1336878
W48762	1336911
W48827	1336957
W49563	1337820
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W49619	1338087
W49672	1337927
W51795	1349846
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W52208	1349338
W52341	1349531
W52355	1349507
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W56077	1357967
W56388	1358278
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	1358409
W56523	
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W56760	1358626
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W58209	1364922
W58291	1365003
W58562	1365294
W58683	1365415
W58731	1365481
W60015	1366774
W60040	1366868
W60288	1367047
W60414	1367398
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W60845	1367603
W60894	1367779
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W68028	1376897
W68141	1377226
W68162	1377032
W68220	1377158
W68291	1377389
W68396	1377267
W69160	1378441
W69213	1378473
W69271	1378746
W69460	1378722
W69649	1379218
W69741	1379072
W69806	1379086
W69960	1379240
W70074	1379335
W70128	1379614
W70147	1379408
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W70242	1379511
W70264	1379553
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W73290	1383425
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W74216	1384396
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W74377	1384792
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W74471	1384758
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W74725	1384948
W74802	1385053
W76133	1386357
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W76313	1386575
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W78967	1389196
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W79340	1390472
W79445	1390696
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W79525	1390785
W79647	1390075
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W79717	1390125
W79779	1390207
W79834	1390242
W80447	1391503
W80496	1391611
W80611	1391698
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W80730	1391748
W80741	1391759
W80791	1391809
W81084	1392136
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W85878	1398307
W85881	1398310
W85931	1398520
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W86124	1398554
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W87865	1401930
W87891	1401976
W88623	1404173
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W88737	1404219
W88786	1404477
W88792	1404264
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W88856	1404408
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X02761	31396
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X04408	31914 35447
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X06617	36143
X06747 X06834	36101 5941
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X07876	33970
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X12791	36112
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X15653	37598
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X16896	33800 36502
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X83973	639692
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X92845	2344811
X92972	1945270
X93036	1085025
X94323	1213612
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X95240	1262818
X95325	1167837
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X97675	1834512
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